

Graph Denoising for Molecular Imaging

Cédric Mendelin <cedric.mendelin@stud.unibas.ch>

Department of Mathematics and Computer Science, University of Basel

29.06.2022

Outline

1 Molecular Imaging Methods

2 Graphs & Manifolds

3 GAT-Denoiser

4 Results

5 Summary & Future Work

6 Questions

Outline

1 Molecular Imaging Methods

2 Graphs & Manifolds

3 GAT-Denoiser

4 Results

5 Summary & Future Work

6 Questions

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

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

Only single particle cryo-EM is considered.

Cryo-Electron Microscopy (Cryo-EM) - Illustration

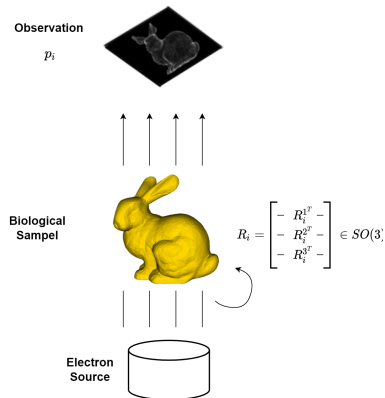


Figure: Cryo-EM overview

Cryo-Electron Microscopy (Cryo-EM) - Illustration

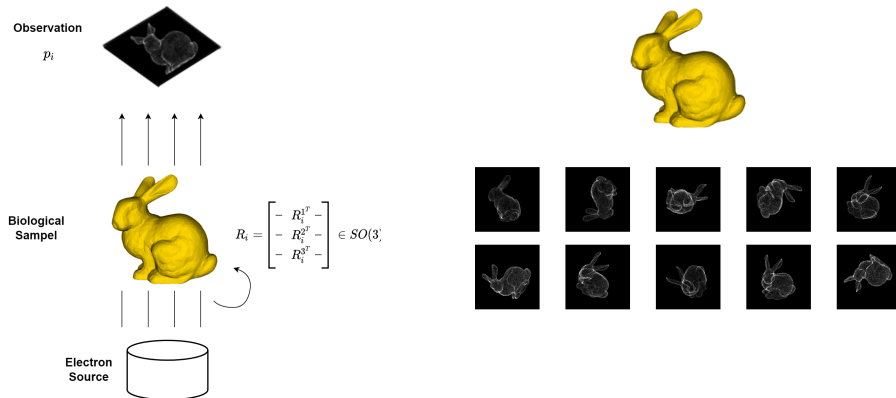
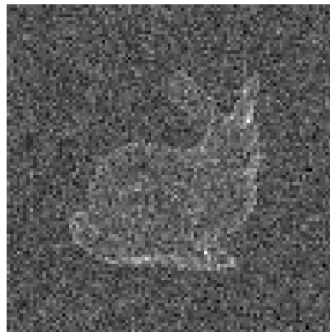


Figure: Cryo-EM overview

Cryo-Electron Microscopy (Cryo-EM) - Illustration



(a) Clean micrograph



(b) Noisy micrograph

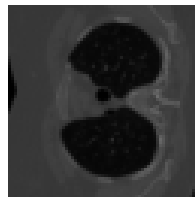
Computed Tomography (CT)

Computed Tomography (CT)

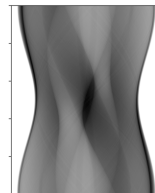
- › Similar to Cryo-EM
- › Can be seen as a simpler version in 2D with known observation angles
- › Good to start with towards a Cryo-EM algorithm

Computed Tomography (CT)

- Similar to Cryo-EM
- Can be seen as a simpler version in 2D with known observation angles
- Good to start with towards a Cryo-EM algorithm



(a) Biological sample



(b) Clean observation (sinogram)

Shared Observation Model

Shared Observation Model

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

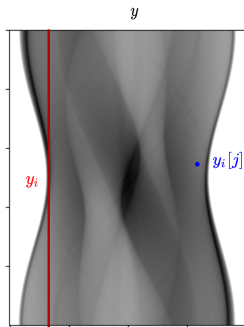
Shared Observation Model

Observation

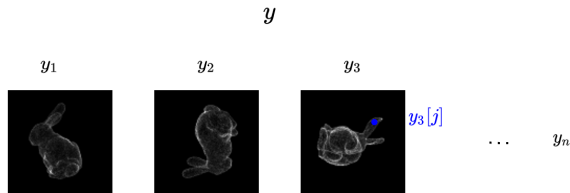
$$\begin{aligned} y_i &= p_i + \eta_i && \text{with } 1 \leq i \leq N \\ y_i &= A(x, \theta_i) + \eta_i && \text{with } 1 \leq i \leq N \end{aligned} \tag{1}$$

- > y : noisy observation
- > p : noiseless observation
- > 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

Reconstruction

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

Signal-to-noise-ratio (SNR)

Signal-to-noise-ratio (SNR)

- SNR is a measure, which compares the power of an input signal to the power of the undesired noise
- Typically given in decibel (dB)
- $\text{SNR} \leq 0$ dB indicated more noise than signal.

Signal-to-noise-ratio (SNR)

- SNR is a measure, which compares the power of an input signal to the power of the undesired noise
- Typically given in decibel (dB)
- $\text{SNR} \leq 0$ dB indicated more noise than signal.

SNR_y is used to define the level of noise in an observation.

Signal-to-noise-ration (SNR)

- SNR is a measure, which compares the power of an input signal to the power of the undesired noise
- Typically given in decibel (dB)
- $\text{SNR} \leq 0$ dB indicated more noise than signal.

SNR_y is used to define the level of noise in an observation.

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

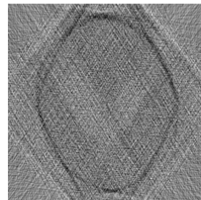
Reconstruction - Computed Tomography

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



(a) Reconstruction clean:

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



(b) Reconstruction noisy

with SNR_y 0 dB:

$$\text{Recon}(y, \theta) \not\approx x$$

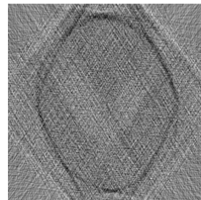
Reconstruction - Computed Tomography

- 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 0 dB:

$$\text{Recon}(y, \theta) \not\approx x$$

Problem and Goal

Problem and Goal

Problem

p not observable directly only access to y .

Problem and Goal

Problem

p not observable directly only access to y .

Goal

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

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

Outline

1 Molecular Imaging Methods

2 Graphs & Manifolds

3 GAT-Denoiser

4 Results

5 Summary & Future Work

6 Questions

Graph - Definitions

Graph - Definitions

Graph Definition

A graph is defined as $G = \langle V, E \rangle$, where V is a set of nodes and E is a set of edges.

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.

Graph - Definitions

Graph Definition

A graph is defined as $G = \langle V, E \rangle$, where V is a set of nodes and E is a set of edges.

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.

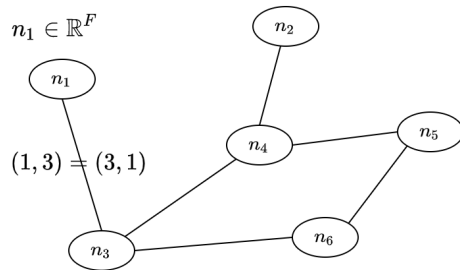


Figure: Sample graph

Graph - Definitions - Adjacency Matrix

Graph - Definitions - Adjacency Matrix

Adjacency Matrix

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

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

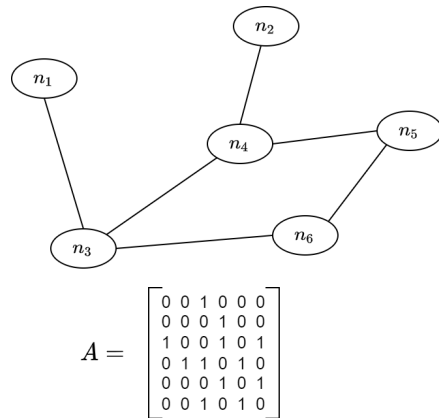


Figure: Sample graph

Graph - Definitions - Degree

Graph - Definitions - Degree

Degree of a node

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

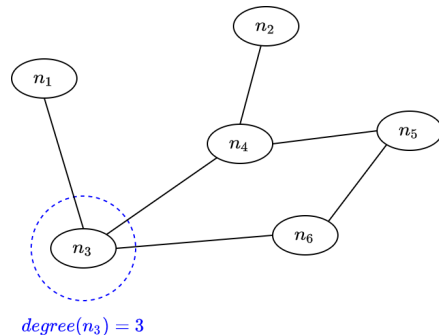


Figure: Sample graph

Graph - Definitions - Degree

Degree of a node

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

Degree Matrix of Graph G

Is a diagonal matrix with degree of nodes as entries.

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

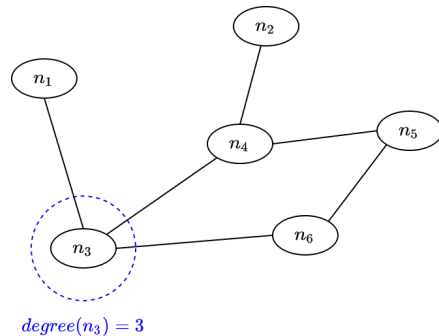


Figure: Sample graph

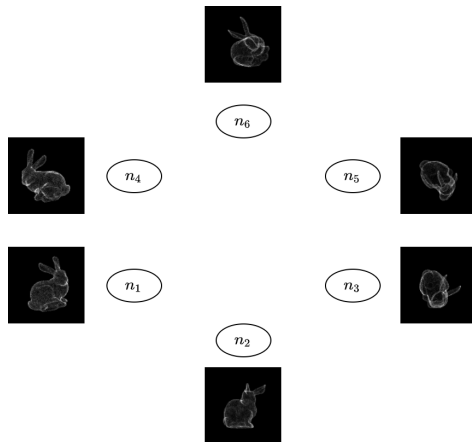
Graph for Molecular Imaging Observation

How to construct a graph for molecular imaging?

Graph for Molecular Imaging Observation

How to construct a graph for molecular imaging?

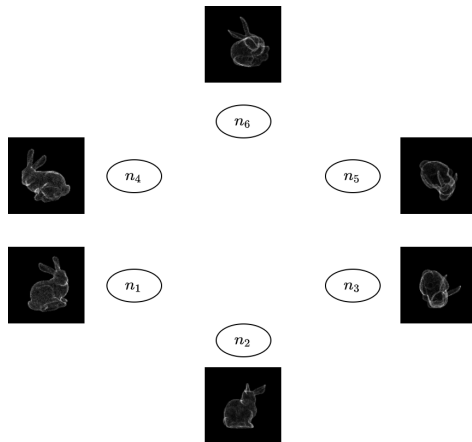
> Nodes: Single observation y_i



Graph for Molecular Imaging Observation

How to construct a graph for molecular imaging?

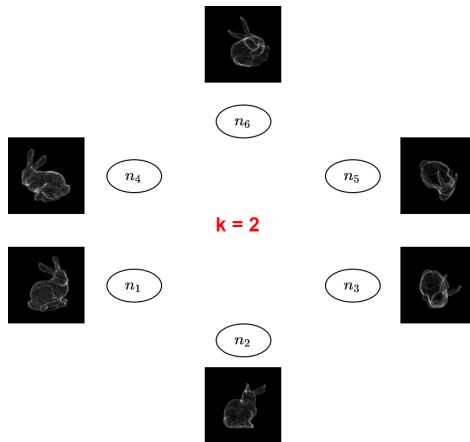
- > Nodes: Single observation y_i
- > Edges: Use k-nearest neighbours (k-NN) to construct a graph



Graph for Molecular Imaging Observation

How to construct a graph for molecular imaging?

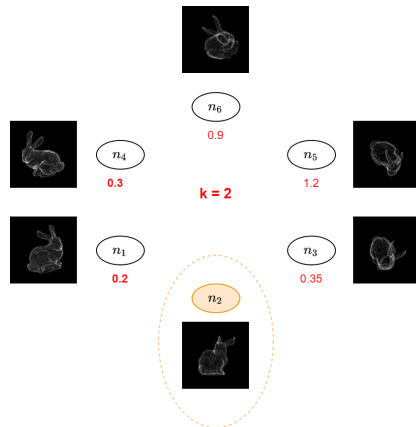
- > Nodes: Single observation y_i
- > Edges: Use k-nearest neighbours (k-NN) to construct a graph
 - > Define similarity measure for nodes: ℓ_2 -norm



Graph for Molecular Imaging Observation

How to construct a graph for molecular imaging?

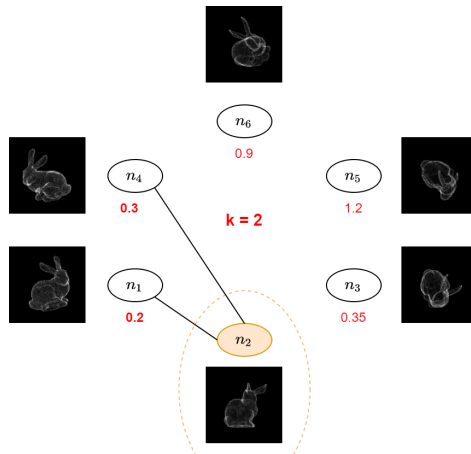
- Nodes: Single observation y_i
- Edges: Use k-nearest neighbours (k-NN) to construct a graph
 - Define similarity measure for nodes: ℓ_2 -norm



Graph for Molecular Imaging Observation

How to construct a graph for molecular imaging?

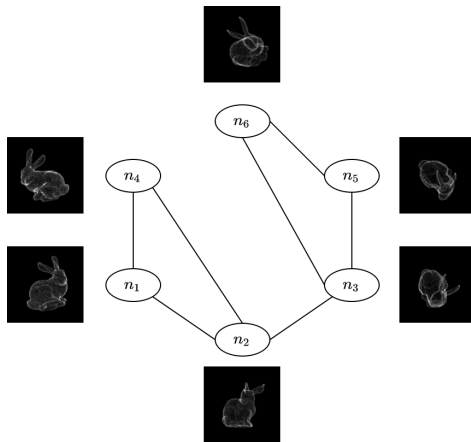
- > Nodes: Single observation y_i
- > Edges: Use k-nearest neighbours (k-NN) to construct a graph
 - > Define similarity measure for nodes: ℓ_2 -norm



Graph for Molecular Imaging Observation

How to construct a graph for molecular imaging?

- › Nodes: Single observation y_i
- › Edges: Use k-nearest neighbours (k-NN) to construct a graph
 - › Define similarity measure for nodes: ℓ_2 -norm



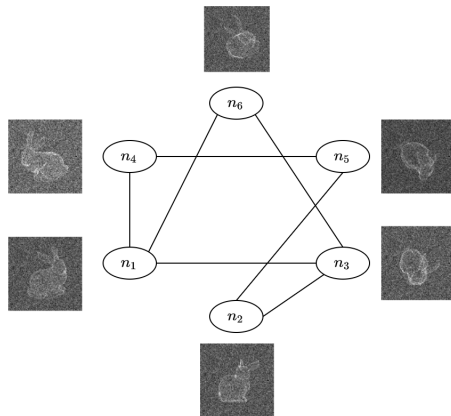
Graph for Molecular Imaging Observation - Noise

What happens with our noisy observations?

Graph for Molecular Imaging Observation - Noise

What happens with our noisy observations?

- > With noise, graph will capture neighborhood inaccurately.



Graph Laplacian (GL)

What can we use this graph for?

Graph Laplacian (GL)

What can we use this graph for?

- Coifman et al. 2008 used it to approximate angles for CT:

Graph Laplacian (GL)

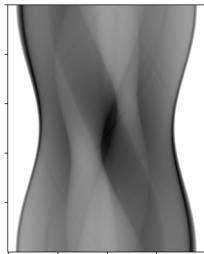
What can we use this graph for?

- Coifman et al. 2008 used it to approximate angles for CT:

Low-dimensional Embedding

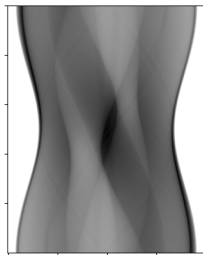
1. Construct a k-NN graph from observations.
2. Calculate $L = D - A$
3. Get 2nd and 3rd smallest eigenvalue with corresponding eigenvectors.

Low-dimensional Embedding for Computed Tomography

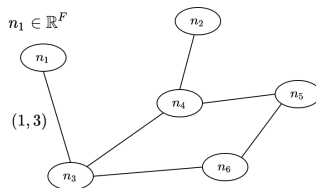


(a) Clean CT observation

Low-dimensional Embedding for Computed Tomography

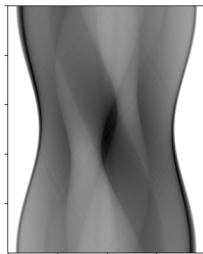


(a) Clean CT observation

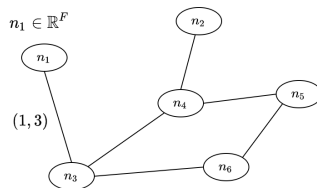


(b) Building k-NN graph with $k = 2$

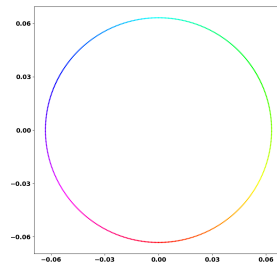
Low-dimensional Embedding for Computed Tomography



(a) Clean CT observation



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



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

Computed Tomography with unknown angles

- GL-Embedding estimates observation angles

Computed Tomography with unknown angles

- GL-Embedding estimates observation angles



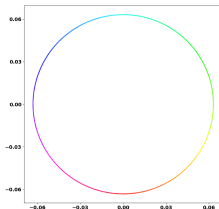
(a) Reconstruction known angles

Computed Tomography with unknown angles

- GL-Embedding estimates observation angles



(a) Reconstruction known angles



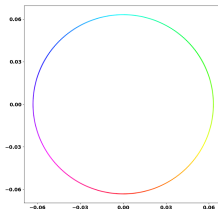
(b) GL-Embedding from $k = 2$

Computed Tomography with unknown angles

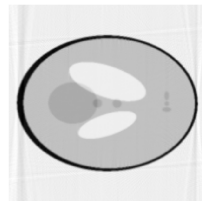
- GL-Embedding estimates observation angles



(a) Reconstruction known angles



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



(c) Reconstruction unknown angles

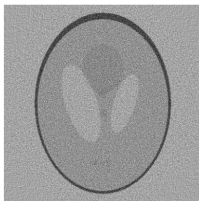
Computed Tomography with unknown angles

- GL-Embedding estimates observation angles

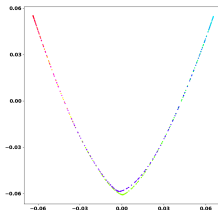
What happens in the noisy case?

Computed Tomography with unknown angles

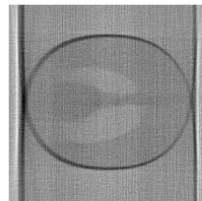
- GL-Embedding estimates observation angles



(a) Reconstruction known angles SNR_y : 10 dB



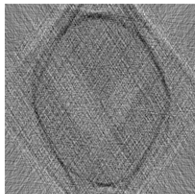
(b) GL-Embedding from $k = 6$ and SNR_y : 10 dB



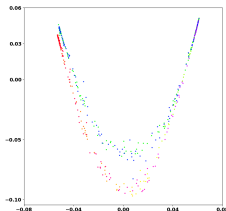
(c) Reconstruction unknown angles SNR_y : 10 dB

Computed Tomography with unknown angles

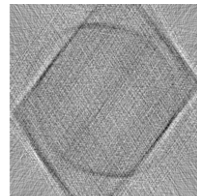
- GL-Embedding estimates observation angles



(a) Reconstruction known angles $SNR_y : 0$ dB



(b) GL-Embedding from $k = 6$ and $SNR_y : 0$ dB



(c) Reconstruction unknown angles $SNR_y : 0$ dB

Observation Denoising

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

Observation Denoising

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

- Use existing denoising algorithms
 - Block-matching and 3D filtering (BM3D) Dabov et al. 2007
 - Non-local means Buades, Coll, and Morel 2005
 - No graph as data structure
 - But, both exploit neighborhood during averaging

Observation Denoising

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

- Use existing denoising algorithms
 - Block-matching and 3D filtering (BM3D) Dabov et al. 2007
 - Non-local means Buades, Coll, and Morel 2005
 - No graph as data structure
 - But, both exploit neighborhood during averaging
- Shows potential for graph as a data structure

Observation Denoising

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

- › Use existing denoising algorithms
 - › Block-matching and 3D filtering (BM3D) Dabov et al. 2007
 - › Non-local means Buades, Coll, and Morel 2005
 - › No graph as data structure
 - › But, both exploit neighborhood during averaging
- › Shows potential for graph as a data structure

Exploit graph as a data structure and the GL-Embedding

Outline

1 Molecular Imaging Methods

2 Graphs & Manifolds

3 GAT-Denoiser

4 Results

5 Summary & Future Work

6 Questions

GAT-Denoiser

- 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

GAT-Denoiser

- › 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

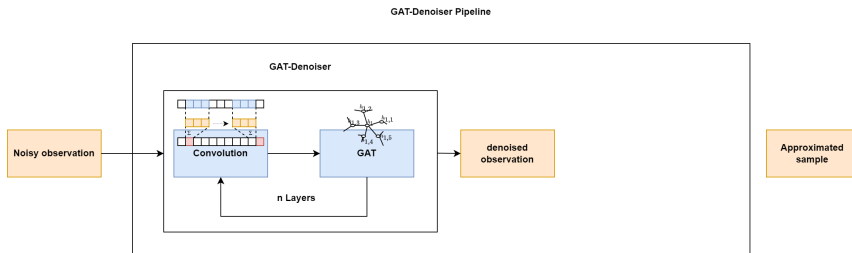
GAT-Denoiser Pipeline

Noisy observation

Approximated
sample

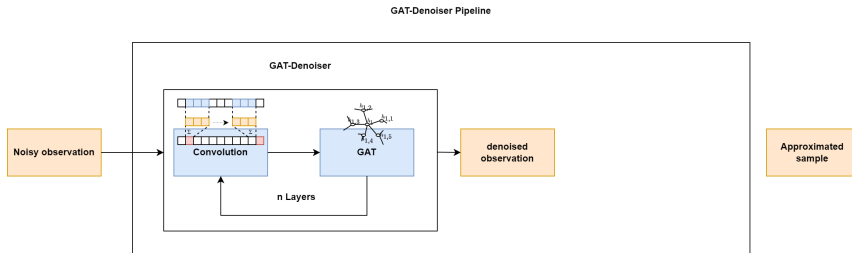
GAT-Denoiser

- 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



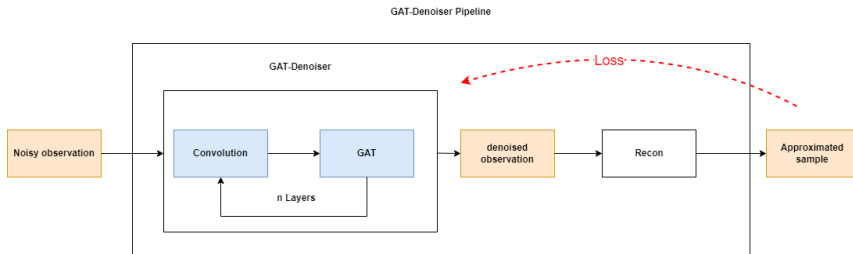
GAT-Denoiser

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



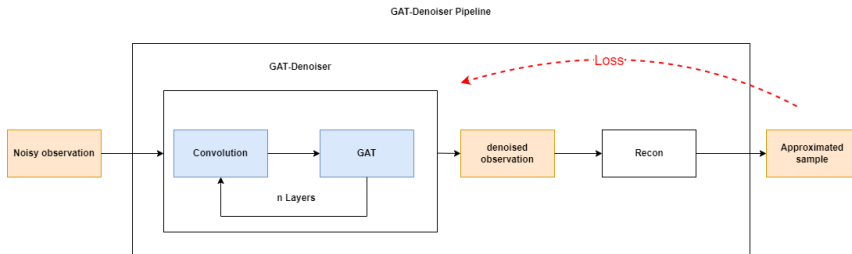
GAT-Denoiser

- > 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**



GAT-Denoiser

- > 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
 - > Optimize for reconstruction quality
 - > Loss: $\mathcal{L}_{reconstruction} = \|x - Recon(GAT-Denoiser(y))\|_2^2$



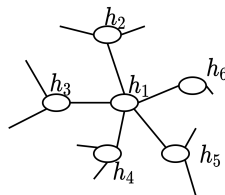
Graph Attention Network - GAT

Graph Attention Network - GAT

- › Extends Graph Convolution Network with attention (weights)
- › Computes new node features
- › Averages graph over neighborhood
- › Multi-head available, motivated by Vaswani et al. 2017

Graph Attention Network - GAT

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



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

Input Graph

Input Graph

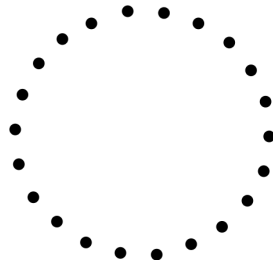
- › Exploit information from GL
 - › Low-dimensional embedding estimates angles
 - › Dominant information in data can be considered observation angles.

Input Graph

- › Exploit information from GL
 - › Low-dimensional embedding estimates angles
 - › Dominant information in data can be considered observation angles.
- › Construct graph from observation angles

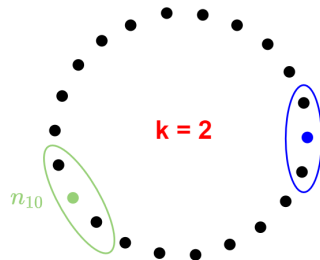
Input Graph

- Exploit information from GL
 - Low-dimensional embedding estimates angles
 - Dominant information in data can be considered observation angles.
- Construct graph from observation angles
 - Map angles to unit-circle / unit-sphere



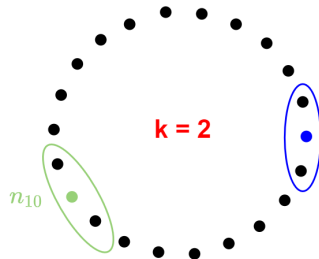
Input Graph

- Exploit information from GL
 - Low-dimensional embedding estimates angles
 - Dominant information in data can be considered observation angles.
- Construct graph from observation angles
 - Map angles to unit-circle / unit-sphere
 - Apply k-NN with great-circle distance



Input Graph

- Exploit information from GL
 - Low-dimensional embedding estimates angles
 - Dominant information in data can be considered observation angles.
- Construct graph from observation angles
 - Map angles to unit-circle / unit-sphere
 - Apply k-NN with great-circle distance



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

Outline

1 Molecular Imaging Methods

2 Graphs & Manifolds

3 GAT-Denoiser

4 **Results**

5 Summary & Future Work

6 Questions

LoDoPaB-CT dataset

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

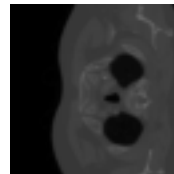
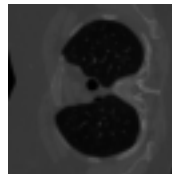
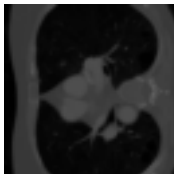


Figure: Some samples from the LoDoPaB-CT dataset.

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
- › Mini-batch gradient descent with batch size 64
- › Adam optimizer

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

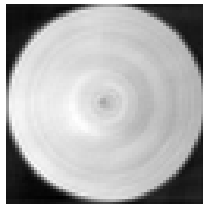
- > 1024 train samples
- > 100 test samples
- > 200 epochs
- > Goal: Find most promising architecture

> Large Scale Experiments

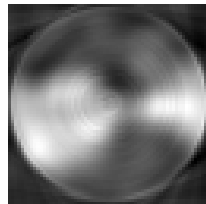
- > Complete LoDoPaB-CT dataset
- > 20 - 40 epochs
- > Goal: Find best model

Small Scale Experiments - Overall Results

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



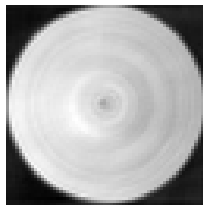
(a) Reconstruction with random Erdős–Rényi graph with $p = 0.01$



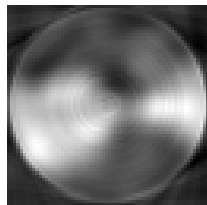
(b) Reconstruction with k-NN input graph $k = 10$

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:
 - › 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 $p = 0.01$

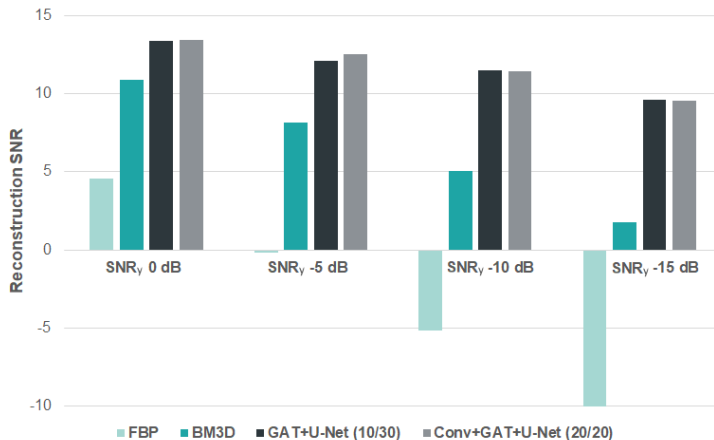


(b) Reconstruction with k-NN input graph $k = 10$

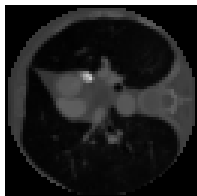
Large Scale Experiment - SNR Results

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

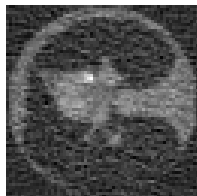
Large Scale Experiment - SNR Results



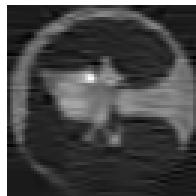
Large Scale Experiment - Visual Results - SNR_y 0 dB



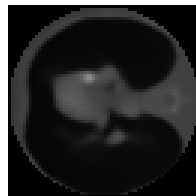
(a) Clean sample



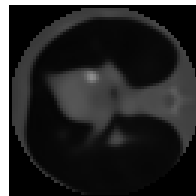
(b) FBP



(c) BM3D



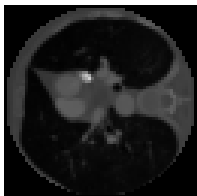
(d) $GAT+U-$
 $Net(10/30)$



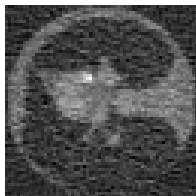
(e)
 $Conv+GAT+U-$
 $Net(20/20)$

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

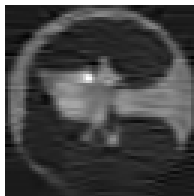
Large Scale Experiment - Visual Results - SNR_y 0 dB



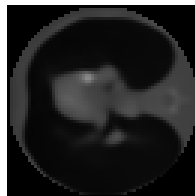
(a) Clean sample



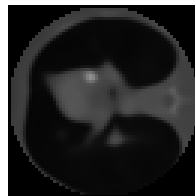
(b) FBP



(c) BM3D



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

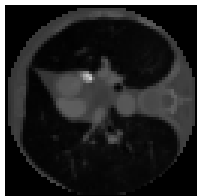


(e)
 $Conv+GAT+U-$
 $Net(20/20)$

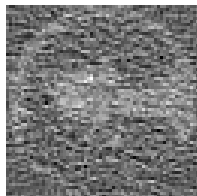
Figure: Large Scale Experiment: Visual results for SNR_y 0 dB.

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

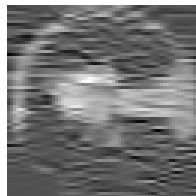
Large Scale Experiment - Visual Results - SNR_y -10 dB



(a) Clean sample



(b) FBP



(c) BM3D



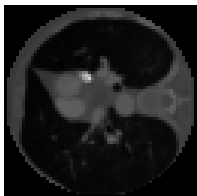
(d) $GAT+U-$
 $Net(10/30)$



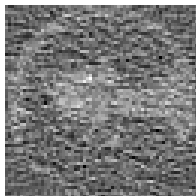
(e)
 $Conv+GAT+U-$
 $Net(20/20)$

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

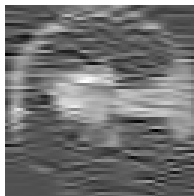
Large Scale Experiment - Visual Results - SNR_y -10 dB



(a) Clean sample



(b) FBP



(c) BM3D



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

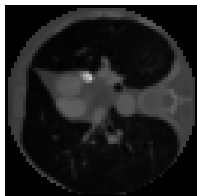


(e)
 $Conv+GAT+U-$
 $Net(20/20)$

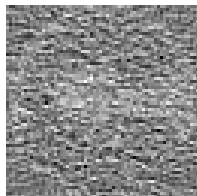
Figure: Large Scale Experiment: Visual results for SNR_y -10 dB.

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

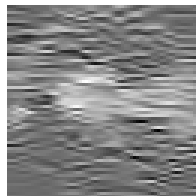
Large Scale Experiment - Visual Results - SNR_y -15 dB



(a) Clean sample



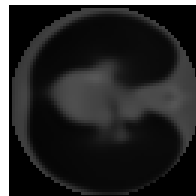
(b) FBP



(c) BM3D



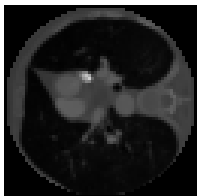
(d) $GAT+U-$
 $Net(10/30)$



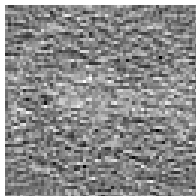
(e)
 $Conv+GAT+U-$
 $Net(20/20)$

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

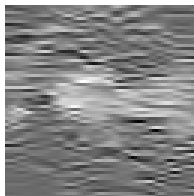
Large Scale Experiment - Visual Results - SNR_y -15 dB



(a) Clean sample



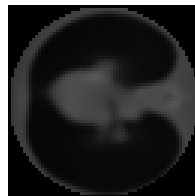
(b) FBP



(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 & Manifolds

3 GAT-Denoiser

4 Results

5 Summary & Future Work

6 Questions

Summary

- › GAT-Denoiser enables denoising of observations
 - › Convolution
 - › GAT
 - › End-To-End Learning
 - › Joint U-Net training boost performance

Summary

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

Future Work

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

Future Work

- › 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

- 1 Molecular Imaging Methods
- 2 Graphs & Manifolds
- 3 GAT-Denoiser
- 4 Results
- 5 Summary & Future Work
- 6 Questions

Questions



References (1)

- Buades, Antoni, Bartomeu Coll, and J-M Morel (2005). “A non-local algorithm for image denoising”. In: **2005 IEEE Computer Society Conference on Computer Vision and Pattern Recognition (CVPR’05)**. Vol. 2. IEEE, pp. 60–65. DOI: 10.1109/CVPR.2005.38.
- Coifman, Ronald R et al. (2008). “Graph Laplacian tomography from unknown random projections”. In: **IEEE Transactions on Image Processing** 17.10, pp. 1891–1899. DOI: 10.1109/TIP.2008.2002305.
- Dabov, Kostadin et al. (2007). “Image denoising by sparse 3-D transform-domain collaborative filtering”. In: **IEEE Transactions on Image Processing** 16.8, pp. 2080–2095. DOI: 10.1109/TIP.2007.901238.
- Leuschner, Johannes et al. (2019). “The lodopab-ct dataset: A benchmark dataset for low-dose ct reconstruction methods”. In: **arXiv preprint arXiv:1910.01113**. DOI: 10.1038/s41597-021-00893-z.

References (2)

- Ronneberger, Olaf, Philipp Fischer, and Thomas Brox (2015). “U-net: Convolutional networks for biomedical image segmentation”. In: **International Conference on Medical image computing and computer-assisted intervention**. Springer, pp. 234–241. DOI: 10.1007/978-3-319-24574-4_28.
- Vaswani, Ashish et al. (2017). “Attention is all you need”. In: **Advances in neural information processing systems** 30. DOI: 10.48550/arXiv.1706.03762.
- Veličković, Petar et al. (2017). “Graph attention networks”. In: **arXiv preprint arXiv:1710.10903**. DOI: 10.48550/arXiv.1710.10903.