

Graph Denoising for Molecular Imaging

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Outline

- Molecular Imaging Methods
- Graphs & Manifolds
- 3 GAT-Denoiser
- 4 Results on LoDoPaB-CT dataset
- 5 Summary & Future Work
- 6 Questions

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Cryo-Electron Microscopy (Cryo-EM)

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- > Enables observation of molecules in near atomic resolution

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 - > During freezing, molecules rotate randomly
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Only single particle cryo-EM is considered.

Graphs & Manifolds

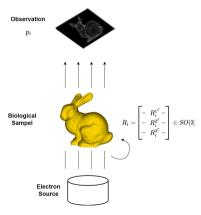


Figure: Cryo-EM overview

Cryo-Electron Microscopy (Cryo-EM) - Illustration

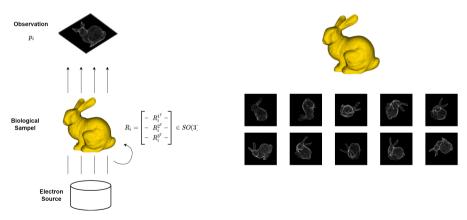


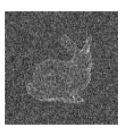
Figure: Cryo-EM overview

Cryo-Electron Microscopy (Cryo-EM) - Challenges

- High-noise level
- > Unknown rotation during freezing
- > (Structural variety of observations)



(a) Clean micrograph



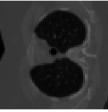
(b) Noisy micrograph

Computed Tomography (CT)

- Related to cryo-EM
- > Can be seen as a simpler version in 2D
- Good to start with towards a cryo-EM algorithm

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(a) Biological sample (b)



(b) Clean observation (sinogram)

Observation

$$y = p + \eta \tag{1}$$

- > y: noisy observation
- > p: noiseless observation
- $>\eta$: noise, assumed $\eta_i \sim \mathcal{N}(0,\sigma^2)$

Observation

$$y = p + \eta$$

$$y_i[j] = p_i[j] + \eta_i[j] \quad \text{with } 1 \le i \le N, 1 \le j \le M$$
(1)

- > v: noisy observation
- p: noiseless observation
- η : noise, assumed $\eta_i \sim \mathcal{N}(0, \sigma^2)$

- > N: number of observations > M: observation dimension

Graphs & Manifolds

Observation

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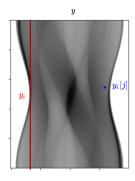
$$y_i[j] = p_i[j] + \eta_i[j] \quad \text{with } 1 \le i \le N, 1 \le j \le M$$

$$y_i = A(x, \theta_i) + \eta_i$$
(1)

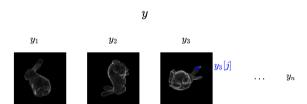
- > v: noisy observation
- p: noiseless observation
- η : noise, assumed $\eta_i \sim \mathcal{N}(0, \sigma^2)$
- > x: biological sample

- > N· number of observations
- > M: observation dimension
- $A: x \mapsto A(x; \theta_i) \in \mathbb{R}^M$: a non-linear operator
- θ_i : observation angle

Observation - Illustration



(a) CT Observation - sinogram



(b) Cryo-EM Observation - micrographs

Reconstruction

Reconstruction

Reconstruction

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

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

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 SNR_{y} is used to define the level of noise in an observation.

SNR is used as a metric for the quality of reconstructions.

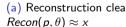
Reconstruction - Computed Tomography

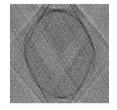
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 - Can be considered historical approach
 - Enables reconstruction for moderate noise

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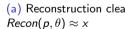
(a) Reconstruction clean: (b) Reconstruction noisy with SNR_v 0 dB: $Recon(y, \theta) \not\approx x$

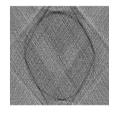
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Graphs & Manifolds

- 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







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Problem and Goal

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p not observable directly only y is observable.

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Goal

denoiser :
$$y_i = (p_i + \eta) \mapsto p_i^* \approx p_i$$

 $Recon(denoiser(y; \theta)) \approx x$

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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.

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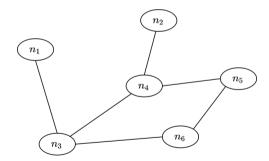


Figure: Sample graph

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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.

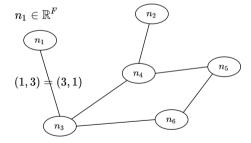


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Graph - Definitions - Adjacency Matrix

Adjacency Matrix

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

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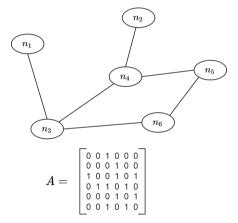


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The *degree* of a node is defined as the number of (incoming) edges.

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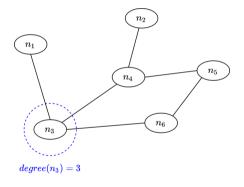


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Degree of a node

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Degree Matrix of Graph G

Is a diagonal matrix with degree of nodes as entries.

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

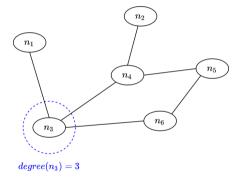


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How to construct a graph for molecular imaging?

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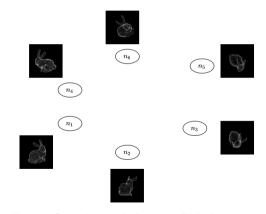


Figure: Sample graph for cryo-EM observation

How to construct a graph for molecular imaging?

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

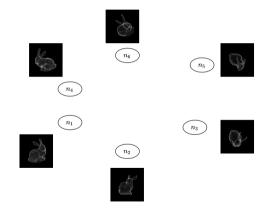


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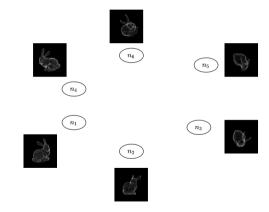


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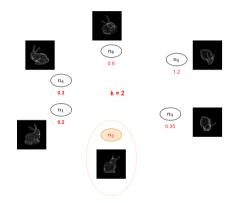


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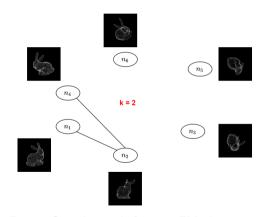


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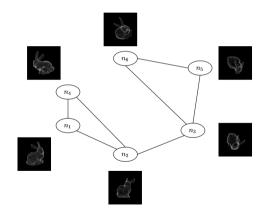


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What happens with our noisy observations?

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With noise, graph will capture neighborhood inaccurately.

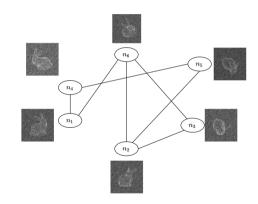


Figure: Sample graph for noisy cryo-EM observation

Graph Laplacian (GL)

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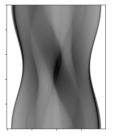
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Low-dimensional Embedding

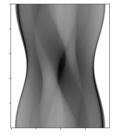
- 1. Construct a k-NN graph from observations.
- 2. Calculate L = D A
- 3. Get 2nd and 3rd smallest eigenvalue with corresponding eigenvectors.

Low-dimensional Embedding for Computed Tomography



(a) Clean CT observation

Low-dimensional Embedding for Computed Tomography

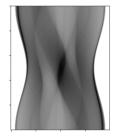


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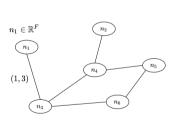


(b) Building k-NN graph with k = 2

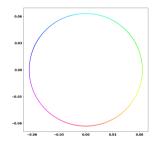
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(c) 2_{nd} and 3_{rd} smallest eigenvectors of L = D - A

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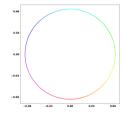


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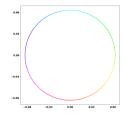


(b) GL-Embedding from k = 2

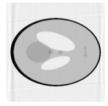
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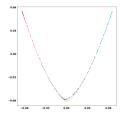
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What happens in the noisy case?

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(a) Reconstruction known angles SNR_v : 10 dB



(b) GL-Embedding from k = 6 and $SNR_v : 10 \text{ dB}$

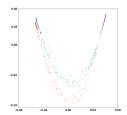


(c) Reconstruction unknown angles SNR_v : 10 dB

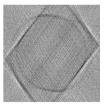
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(a) Reconstruction known angles SNR_v : 0 dB



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Observation Denoising

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 - Non-local means Buades, Coll, and Morel 2005
 - No graph as data structure
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olecular Imaging Methods Graphs & Manifolds GAT-Denoiser Results on LoDoPaB-CT dataset Summary & Future Work Ques

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Exploit graph as a data structure and the GL-embedding.

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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

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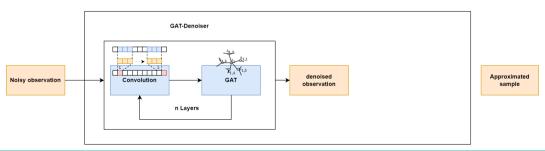


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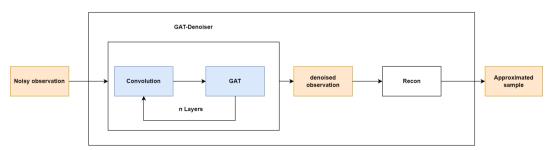


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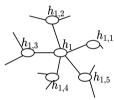
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- End-to-End Learning
 - Optimize for reconstruction quality
 - $\mathcal{L} = ||x Recon(GAT-Denoiser(A(x, \theta) + \eta))||_2^2$
 - $\mathcal{L}_{sino} = \| p GAT Denoiser(A(x, \theta) + n) \|_2^2$

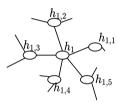
Graphs & Manifolds

- Extends Graph Convolution Network with attention (weights)
- Compute new node features
- Averages graph over neighborhood
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Graphs & Manifolds

- Extends Graph Convolution Network with attention (weights)
- Compute 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
- $\geq \alpha$: normalized attention coefficients



$$h_1' = \sigma \left(\sum_{i=1}^5 \alpha_i W h_{1,i} \right)$$

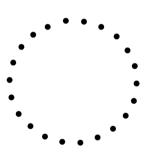
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- > Low-dimensional embedding estimates angles
- Dominant information in data can be considered observation angles.

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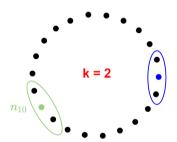
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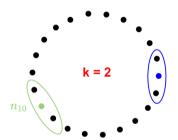


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GAT-Denoiser Implementation for Computed Tomography

- Use U-Net for reconstruction
- > During Trainig, U-Net might be trained jointly

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LoDoPaB-CT dataset

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









Figure: Some samples from the LoDoPaB-CT dataset.

Evaluation

- > Small Scale Experiments
 - > 1024 train samples
 - > 100 test samples
 - > 200 epochs
- Large Scale Experiments
 - > Complete LoDoPaB-CT dataset
 - > 20 40 epochs

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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

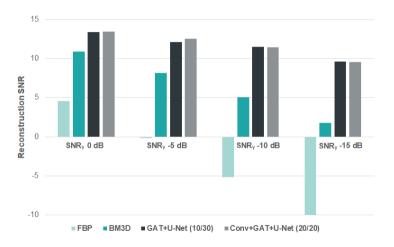
Training

- U-Net used for reconstruction
 - \rightarrow Pre-trained with complete dataset and SNR_y in [-10, 0] for 200 epochs
- > Mini-batch gradient descent with batch size 64
- > Adam optimizer
- > Joint U-Net training possible

Small Scale Results

- > Learning fails with random graph
- Learning succeeds with defined input graph
- Components contribute to success of GAT-Denoiser
- > Best model with joint U-Net training

Large Scale Results



Large Scale Results - Visual - SNR_v 0 dB

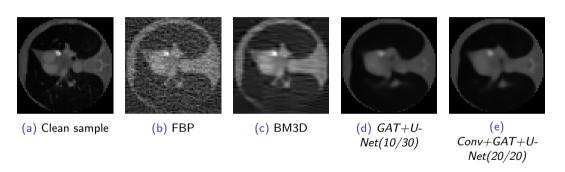


Figure: Large Scale Experiment: Visual results for SNR_v 0 dB.

Large Scale Results - Visual - SNR_v 0 dB

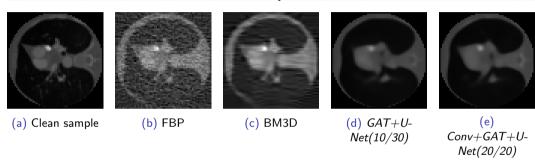


Figure: Large Scale Experiment: Visual results for SNR_v 0 dB.

GAT-Denoiser improves BM3D by 27.6%.

Large Scale Results - Visual - SNR_v -10 dB

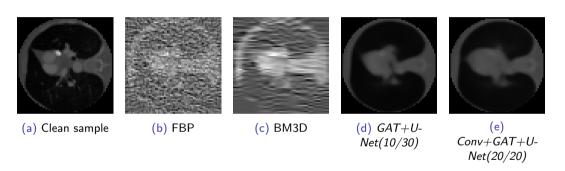


Figure: Large Scale Experiment: Visual results for SNR_v -10 dB.

Large Scale Results - Visual - SNR_v -10 dB

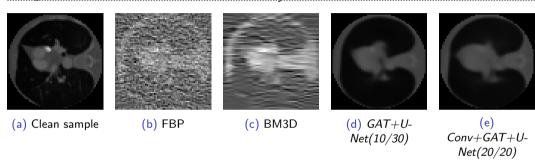


Figure: Large Scale Experiment: Visual results for SNR_v -10 dB.

GAT-Denoiser improves BM3D by 126.0%.

Large Scale Results - Visual - SNR_v -15 dB

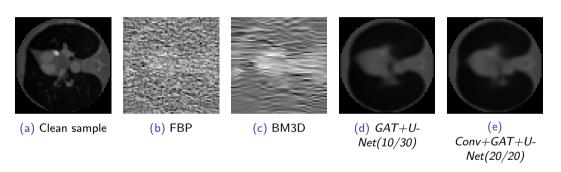


Figure: Large Scale Experiment: Visual results for SNR_v -15 dB.

Large Scale Results - Visual - SNR_v -15 dB

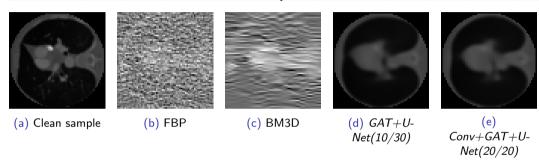


Figure: Large Scale Experiment: Visual results for SNR_v -15 dB.

GAT-Denoiser improves BM3D by 379.9%.

Outline

- Molecular Imaging Methods
- Graphs & Manifolds
- GAT-Denoiser
- 4 Results on LoDoPaB-CT datases
- 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

olecular Imaging Methods Graphs & Manifolds GAT-Denoiser Results on LoDoPaB-CT dataset **Summary & Future Work** Ques

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

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- 6 Questions

Questions

Molecular Imaging Methods Graphs & Manifolds GAT-Denoiser Results on LoDoPaB-CT dataset Summary & Future Work Questions

Questions



References

- Basu, Samit and Yoram Bresler (2000). "Feasibility of tomography with unknown view angles". In: **IEEE Transactions on Image Processing 9.6, pp. 1107–1122.** DOI: 10.1109/83.846252.
- Bendory, Tamir, Alberto Bartesaghi, and Amit Singer (2020). "Single-particle cryo-electron microscopy: Mathematical theory, computational challenges, and opportunities". In: IEEE Signal Processing Magazine 37.2, pp. 58–76. DOI: 10.1109/MSP.2019.2957822.
- Biewald, Lukas (2020). Experiment Tracking with Weights and Biases. Software available from wandb.com. URL: https://www.wandb.com/.
- Brenner, David J and Eric J Hall (2007). "Computed tomography—an increasing source of radiation exposure". In: New England journal of medicine 357.22, pp. 2277–2284. DOI: 10.1056/NEJMra072149.
- 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.
- Cayton, Lawrence (2005). "Algorithms for manifold learning". In: Univ. of California at San Diego **Tech. Rep** 12.1-17, p. 1.
- Clackdoyle, Rolf and Michel Defrise (2010). "Tomographic reconstruction in the 21st century". In: