

# Application of Graph Learning to inverse problems

---

Cédric Mendelin <[cedric.mendelin@stud.unibas.ch](mailto:cedric.mendelin@stud.unibas.ch)>

Department of Mathematics and Computer Science, University of Basel

Date

# Outline

---

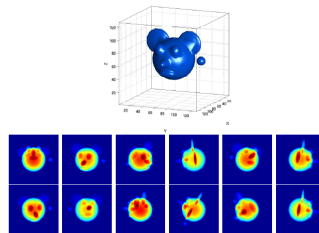
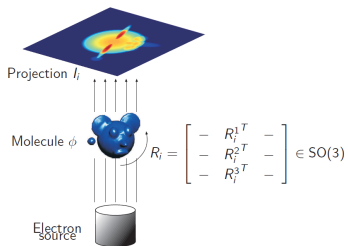
1 Imaging methods

2 Graph Denoising

3 Conclusion

# Cryo-EM

---



Cryo-EM overview Singer 2018, Figure 1 and Figure 2

## Cryo-EM challenges

---

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

## Cryo-EM challenges

---

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

Only single particle cryo-EM is considered.

## Cryo-EM challenges

---

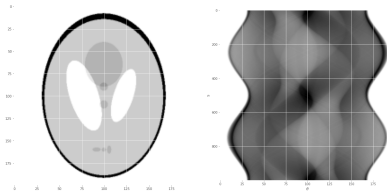
- High-noise level
- Unknown rotation during freezing
- (Structural variety of observations)

Only single particle cryo-EM is considered.

The main domain of interest is high-noise domain (cryo-EM).

# Computed Tomography

---



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

## Graph construction

---

Consider  $n$  observed images  $(x_0, x_1, \dots, x_n)$ , where  $x_i \in \mathbb{R}^M$  with  $M$  as image dimension. A graph can be constructed by:

- $V$ : The images can be used as nodes.
- $E$ : If similarity measure  $d$  of two images  $x_i, x_j$  is smaller than given threshold  $\tau$ , there will be an edge.



## Graph construction

---

Consider  $n$  observed images  $(x_0, x_1, \dots, x_n)$ , where  $x_i \in \mathbb{R}^M$  with  $M$  as image dimension. A graph can be constructed by:

- $V$ : The images can be used as nodes.
- $E$ : If similarity measure  $d$  of two images  $x_i, x_j$  is smaller than given threshold  $\tau$ , there will be an edge.

### Definition (Adjacency Matrix)

$$A_{ij} = \begin{cases} 1 & \text{if } d(x_i, x_j) < \tau \\ 0, & \text{otherwise} \end{cases} \quad (1)$$

## Graph construction

---

Consider  $n$  observed **noisy** images  $(y_0, y_1, \dots, y_n)$ , where  $y_i \in \mathbb{R}^M$  with  $M$  as image dimension. Then,  $y_i = x_i + \eta$  with  $\eta$  drawn from  $\mathcal{N} \sim (0, \sigma^2)$  and  $(x_0, x_1, \dots, x_n)$  as the original images. A noisy graph can be constructed like:

- $V$ : The **noisy** images  $y$  can be used as nodes.
- $E$ : If similarity measure  $d$  of two **noisy** images  $y_i, y_j$  is smaller than given threshold  $\tau$ , there will be an edge.

## Graph construction

---

Consider  $n$  observed **noisy** images  $(y_0, y_1, \dots, y_n)$ , where  $y_i \in \mathbb{R}^M$  with  $M$  as image dimension. Then,  $y_i = x_i + \eta$  with  $\eta$  drawn from  $\mathcal{N} \sim (0, \sigma^2)$  and  $(x_0, x_1, \dots, x_n)$  as the original images. A noisy graph can be constructed like:

- $V$ : The **noisy** images  $y$  can be used as nodes.
- $E$ : If similarity measure  $d$  of two **noisy** images  $y_i, y_j$  is smaller than given threshold  $\tau$ , there will be an edge.

### Definition (Noisy Adjacency Matrix)

$$A_{ij} = \begin{cases} 1 & \text{if } d(y_i, y_j) < \tau \\ 0, & \text{otherwise} \end{cases} \quad (2)$$

# Graph Denoising

---

*Graph denoising* is the task, to estimate a denoised graph  $\tilde{G}$  from a given noisy graph  $G_0$ , with underlying original graph  $G$ :

## Definition (Graph Denoising)

$$GD : G_0 \mapsto \tilde{G} \approx G,$$

where  $G_0$ ,  $\tilde{G}$ ,  $G$  denotes noisy, estimated denoised and original graph respectively.

Slide about Manifolds on cryo-EM and CT.

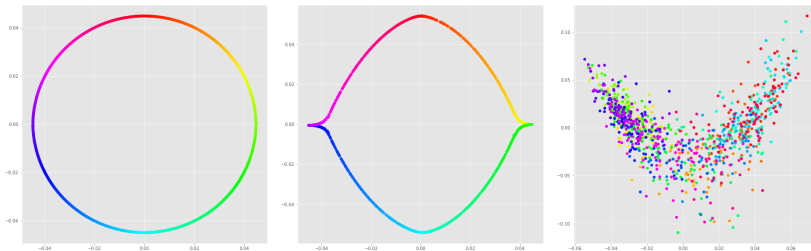
# Manifolds

---

We know structure of manifolds for noiseless data:

- Computed Tomography → circle
- Cryo-Em → sphere

If data is noisy, manifold will "drift" away from noiseless manifold:



Manifolds for phantom without noise, and different noise levels.

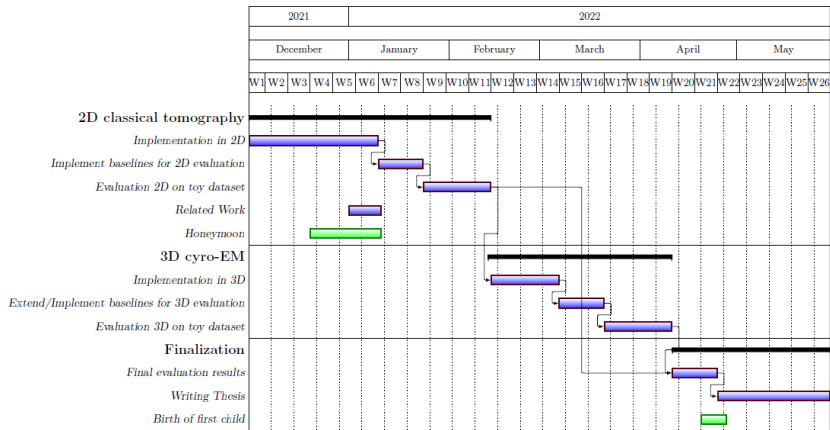
## Project Conclusion - Next steps

---

- Focus on high-noise domain (cryo-EM)
- Introduce Graph Denoising method
- Exploit known manifold in 2D and 3D
- Further study Graph Laplacian and connection with
  - Tomography domain
  - GNNs and Machine Learning in general
- Evaluate on toy dataset in 2D and 3D
- Nice to have: evaluate on cryo-EM / CT dataset

# Project schedule

---





Questions?

# References

---

- Singer, Amit (2018). “Mathematics for cryo-electron microscopy”. In: **Proceedings of the International Congress of Mathematicians: Rio de Janeiro 2018**. World Scientific, pp. 3995–4014.