UNIVERSITY OF COPENHAGEN

Graph Refinement based Airway Extraction Using Mean-field and Graph Neural Networks

Raghavendra Selvan raghav@di.ku.dk Department of Computer Science

Outline

Airway Diseases

2 Understanding the Data

3 Methods

Existing Methods Graph Refinement Model Mean-Field Networks Graph Neural Networks

4 Experiments

Outline

Airway Diseases

Output Description of the Data

Existing Methods Graph Refinement Model Mean-Field Networks Graph Neural Networks

O Experiments



- Tobacco Smoking
- Indoor & Outdoor Air pollution



- Tobacco Smoking
- Indoor & Outdoor Air pollution
- Third leading cause of death by 2030
- 174.5M affected; 3.2M deaths (2015)



- Tobacco Smoking
- Indoor & Outdoor Air pollution
- Third leading cause of death by 2030
- 174.5M affected; 3.2M deaths (2015)
- Destruction of lung tissue (Emphysema)
- Change of airway morphology (Bronchiectasis)



- Tobacco Smoking
- Indoor & Outdoor Air pollution
- Third leading cause of death by 2030
- 174.5M affected; 3.2M deaths (2015)
- Destruction of lung tissue (Emphysema)
- Change of airway morphology (Bronchiectasis)





- Tobacco Smoking
- Indoor & Outdoor Air pollution
- Third leading cause of death by 2030
- 174.5M affected; 3.2M deaths (2015)
- Destruction of lung tissue (Emphysema)
- Change of airway morphology (Bronchiectasis)





Existing Diagnostics are Rudimentary & Tedious

- Lung Function Tests
 - + Simple and inexpensive
 - Patient dependent
 - Low reproducibility
 - Mild cases go unnoticed



Existing Diagnostics are Rudimentary & Tedious

- Lung Function Tests
 - + Simple and inexpensive
 - Patient dependent
 - Low reproducibility
 - Mild cases go unnoticed
- 3D CT Scans
 - + Provide more information
 - Arduous to read the data; even for experts
 - Low inter-observer agreement







Automatic Airway Segmentation and obtain useful COPD biomarkers





Outline

Airway Diseases

Output Data

Methods

Existing Methods Graph Refinement Model Mean-Field Networks Graph Neural Networks





Primary Data from Danish Lung Cancer Study

- Danish Lung Cancer Screening Trial ¹
- Low-dose CT
- $\bullet \ > 10,000 \text{ scans}$
- Age 50-70 years.
- Smoker or former smoker (> 20 pack years)
- 32/10,000 have segmentations verified by an expert user!

¹Pedersen, J. H., et.al : The Danish randomized lung cancer CT screening trialoverall design and results of the prevalence round. Journal of Thoracic Oncology, (2009)



Slide 8 — Raghavendra Selvan — Graph Refinement based Airway Extraction

CT Images are noisy, low contrast & low-res.



- Volume resolution \sim 300x250x275
- Voxels \sim 0.75mmx0.75mmx1mm
- Challenges
 - Acquisition noise
 - Inter-patient variability
 - Several "interfering" structures
 - Labels/Annotations



Outline

Airway Diseases

Output Description of the Data

Methods Existing Methods Graph Refinement Model Mean-Field Networks Graph Neural Networks

Most Existing Methods handle occlusions poorly

- State-of-the-art: Region-growing (!) based methods
- EXACT Study² compares 15 methods; No clear winner
- Small airways are challenging
- Challenging to overcome occlusions

 $^2 \text{Lo},$ P., et.al : Extraction of airways from CT (EXACT'09). IEEE Transactions on Medical Imaging, (2012)



Slide 11 — Raghavendra Selvan — Graph Refinement based Airway Extraction

Graph Refinement Model for Airway Extraction



Graph Refinement Model for Airway Extraction

Desired Properties

- Exploratory (to overcome occlusions)
- Detect small airways
- Uncertainty estimates



Preprocess Image to Graph Model

One possibility



Preprocess Image to Graph Model

One possibility



Figure 1: Visualisation of the pre-processing carried out to transform the input image (left) into a probability image (center) and then into graph format (right). Nodes in the graph are shown in scale to capture the variations in their local radius.



Graph Refinement Model

- Input graph: $G_i : \{N, E_i\}$
- Node features: $\mathbf{X} \in \mathbb{R}^{F \times N}$
- Input adjacency: $\mathbf{A}_i \in \{0,1\}^{N \times N}$



Graph Refinement Model

- Input graph: $G_i : \{N, E_i\}$
- Node features: $\mathbf{X} \in \mathbb{R}^{F \times N}$
- Input adjacency: $\mathbf{A}_i \in \{0,1\}^{N \times N}$

Graph Refinement Task

$$f(\mathcal{G}_i)
ightarrow \mathcal{G}$$

Output subgraph \mathcal{G} with $\mathcal{E} \subset \mathcal{E}_i$; $\mathbf{A} \in \{0, 1\}^{N \times N}$

Visualise Graph Refinement Task





Visualise Graph Refinement Task



- Binary random variable to capture existence of edge between nodes
- $s_{ij} \in \{0,1\}$ with prob. $\alpha_{ij} \in [0,1]$



- Binary random variable to capture existence of edge between nodes
- $s_{ij} \in \{0,1\}$ with prob. $\alpha_{ij} \in [0,1]$
- For each node: $\mathbf{s}_i = \{s_{ij}\} : j = 1 ... N$
- Global connectivity variable: $\boldsymbol{S} = [\boldsymbol{s}_1 \dots \boldsymbol{s}_N]$
- Each instance of **S** an $N \times N$ adjacency matrix



- Binary random variable to capture existence of edge between nodes
- $s_{ij} \in \{0,1\}$ with prob. $\alpha_{ij} \in [0,1]$
- For each node: $\mathbf{s}_i = \{s_{ij}\} : j = 1 ... N$
- Global connectivity variable: $\boldsymbol{S} = [\boldsymbol{s}_1 \dots \boldsymbol{s}_N]$
- Each instance of **S** an $N \times N$ adjacency matrix

Input Graph G_i is completely described by \mathbf{X}, \mathbf{A}_i ,



- Binary random variable to capture existence of edge between nodes
- $s_{ij} \in \{0,1\}$ with prob. $\alpha_{ij} \in [0,1]$
- For each node: $\mathbf{s}_i = \{s_{ij}\} : j = 1 ... N$
- Global connectivity variable: $\boldsymbol{S} = [\boldsymbol{s}_1 \dots \boldsymbol{s}_N]$
- Each instance of **S** an $N \times N$ adjacency matrix

Input Graph G_i is completely described by \mathbf{X}, \mathbf{A}_i ,

Posterior density of interest: $p(\mathbf{S}|\mathbf{X}, \mathbf{A}_i)$



- Binary random variable to capture existence of edge between nodes
- $s_{ij} \in \{0,1\}$ with prob. $\alpha_{ij} \in [0,1]$
- For each node: $\mathbf{s}_i = \{s_{ij}\} : j = 1 ... N$
- Global connectivity variable: $\boldsymbol{S} = [\boldsymbol{s}_1 \dots \boldsymbol{s}_N]$
- Each instance of **S** an $N \times N$ adjacency matrix

Input Graph G_i is completely described by \mathbf{X}, \mathbf{A}_i ,

Posterior density of interest: $p(\mathbf{S}|\mathbf{X}, \mathbf{A}_i)$

$$\ln p(\mathbf{S}|\mathbf{X}) \propto \ln p(\mathbf{S}, \mathbf{X})$$

= $-\ln Z + \sum_{i \in \mathcal{N}} \phi_i(\mathbf{s}_i) + \sum_{(i,j) \in \mathcal{E}} \phi_{ij}(\mathbf{s}_i, \mathbf{s}_j)$





 $p(\mathbf{S}|\mathbf{X}, \mathbf{A}_i)$ is intractable except for in trivial cases.



 $p(\mathbf{S}|\mathbf{X}, \mathbf{A}_i)$ is intractable except for in trivial cases.

 $p(\mathbf{S}|\mathbf{X},\mathbf{A}_i) \approx q(\mathbf{S})$



 $p(\mathbf{S}|\mathbf{X}, \mathbf{A}_i)$ is intractable except for in trivial cases.

 $p(\mathbf{S}|\mathbf{X},\mathbf{A}_i) pprox q(\mathbf{S})$

Variational Approximate Inference

Minimize ELBO to obtain $q(\mathbf{S}) \in \mathcal{Q}$

$$\mathcal{F}(q_{\mathsf{S}}) = \ln Z + \mathbb{E}_{q_{\mathsf{S}}} \Big[\ln p(\mathsf{S}|\mathsf{X}) - \ln q(\mathsf{S}) \Big]$$



Simpler approximation using MFA



Simpler approximation using MFA

Factorisable $q(\mathbf{S})$

$$q(\mathbf{S}) = \prod_{i=1}^{N} \prod_{j=1}^{N} q_{ij}(s_{ij}),$$

where, $q_{ij}(s_{ij}) = \begin{cases} lpha_{ij} & \text{if } s_{ij} = 1\\ (1 - lpha_{ij}) & ext{if } s_{ij} = 0 \end{cases},$



Simpler approximation using MFA

Factorisable
$$q(\mathbf{S})$$

$$q(\mathbf{S}) = \prod_{i=1}^{N} \prod_{j=1}^{N} q_{ij}(s_{ij}),$$
where, $q_{ij}(s_{ij}) = \begin{cases} \alpha_{ij} & \text{if } s_{ij} = 1\\ (1 - \alpha_{ij}) & \text{if } s_{ij} = 0 \end{cases}$

Assumes edges to be independent.



Node and Pairwise Potentials for MFA



Node and Pairwise Potentials for MFA

Node Potential

For each node $i \in \mathcal{N}$,

$$\phi_i(\mathbf{s}_i) = \sum_{\mathbf{v}=0}^D \beta_{\mathbf{v}} \mathbb{I}\Big[\sum_j s_{ij} = \mathbf{v}\Big] + \mathbf{a}^T \mathbf{x}_i \sum_j s_{ij},$$



Node and Pairwise Potentials for MFA

Node Potential

For each node $i \in \mathcal{N}$,

$$\phi_i(\mathbf{s}_i) = \sum_{\mathbf{v}=0}^D \beta_{\mathbf{v}} \mathbb{I}\Big[\sum_j s_{ij} = \mathbf{v}\Big] + \mathbf{a}^T \mathbf{x}_i \sum_j s_{ij},$$

Pairwise Potential

For each edge, $(i, j) \in \mathcal{E}_i$

$$\phi_{ij}(\mathbf{s}_i, \mathbf{s}_j) = \lambda \big(1 - 2|s_{ij} - s_{ji}| \big) + (2s_{ij}s_{ji} - 1) \Big[\boldsymbol{\eta}^T |\mathbf{x}_i - \mathbf{x}_j| + \boldsymbol{\nu}^T (\mathbf{x}_i \mathbf{x}_j) \Big]$$

 $\mathsf{Parameters} = [\boldsymbol{\beta}, \mathbf{a}, \lambda, \boldsymbol{\eta}, \boldsymbol{\nu}]$



Minimize ELBO to get MFA Iterations

MFA Iterations

$$\alpha_{kl}^{(t+1)} = \sigma(\gamma_{kl}) = \frac{1}{1 + \exp^{-\gamma_{kl}}} \forall k = \{1 \dots N\}, \ l \in \mathcal{N}_k,$$

where $\sigma(\cdot)$ is sigmoid activation, \mathcal{N}_k are neighbours of node k, and



Minimize ELBO to get MFA Iterations

MFA Iterations

$$\alpha_{kl}^{(t+1)} = \sigma(\gamma_{kl}) = \frac{1}{1 + \exp^{-\gamma_{kl}}} \forall k = \{1 \dots N\}, \ l \in \mathcal{N}_k,$$

where $\sigma(\cdot)$ is sigmoid activation, \mathcal{N}_k are neighbours of node k, and

$$\gamma_{kl} = \prod_{j \in \mathcal{N}_k \setminus l} \left(1 - \alpha_{kj}^{(t)}\right) \left\{ \sum_{m \in \mathcal{N}_k \setminus l} \frac{\alpha_{km}^{(t)}}{(1 - \alpha_{km}^{(t)})} \left[(\beta_2 - \beta_1) - \beta_2 \sum_{n \in \mathcal{N}_k \setminus l, m} \frac{\alpha_{kn}^{(t)}}{(1 - \alpha_{kn}^{(t)})} \right] + (\beta_1 - \beta_0) \right\} + \mathbf{a}^T \mathbf{x}_k + \left(4\alpha_{lk}^{(t)} - 2)\lambda + 2\alpha_{lk}^{(t)} (\boldsymbol{\eta}^T | \mathbf{x}_k - \mathbf{x}_l | + \boldsymbol{\nu}^T (\mathbf{x}_k \mathbf{x}_l))\right).$$
(1)

1.5

MFA to Mean-Field Networks



MFA to Mean-Field Networks

• MFA Iterations resemble feed-forward operations in neural network

$$\alpha_{kl}^{(t+1)} = \sigma(\gamma_{kl}) = \frac{1}{1 + \exp^{-\gamma_{kl}}} \ \forall \ k = \{1 \dots N\}, \ l \in \mathcal{N}_k,$$

lpha is soft prediction of global connectivity variable



MFA to Mean-Field Networks

• MFA Iterations resemble feed-forward operations in neural network

$$\alpha_{kl}^{(t+1)} = \sigma(\gamma_{kl}) = \frac{1}{1 + \exp^{-\gamma_{kl}}} \forall k = \{1 \dots N\}, \ l \in \mathcal{N}_k,$$

lpha is soft prediction of global connectivity variable

- *T*-iterations as a *T*-layered network
- Automatic differentiation to learn parameters: $\mathcal{L}(\boldsymbol{\alpha}, \mathbf{A}_r)$, where \mathbf{A}_r is reference adjacency.



Summarising MFN



Summarising MFN

- Yields approximation to underlying posterior
- Simple factors
- Handful of parameters
- Easy to optimise
- Hand-crafting potentials is cumbersome
- Might not generalise across applications





 Neural networks operating directly on graph structured data



- Neural networks operating directly on graph structured data
- Generalisation of message passing algorithms



- Neural networks operating directly on graph structured data
- Generalisation of message passing algorithms
- Arbitrarily complex messages



- Neural networks operating directly on graph structured data
- Generalisation of message passing algorithms
- Arbitrarily complex messages
- End-to-end trainable inference systems



- Neural networks operating directly on graph structured data
- Generalisation of message passing algorithms
- Arbitrarily complex messages
- End-to-end trainable inference systems



Figure: Two mappings of Factor Graphs into GNNs

Figure from Yoon et al. "Inference in probabilistic graphical models by Graph Neural Networks" (2018)



• Encoder comprises of GNNs; Message passing between nodes



- Encoder comprises of GNNs; Message passing between nodes
- Jointly train to learn useful embeddings



- Encoder comprises of GNNs; Message passing between nodes
- Jointly train to learn useful embeddings
- Suitable for graph refinement: $f(\mathcal{G}_i) \rightarrow \mathcal{G}$



- Encoder comprises of GNNs; Message passing between nodes
- Jointly train to learn useful embeddings
- Suitable for graph refinement: $f(\mathcal{G}_i) \rightarrow \mathcal{G}$



GAE Model for Graph Refinement: Encoder



GAE Model for Graph Refinement: Encoder

Encoder:

Node Embedding: Node-to-Edge mapping:

Edge-to-Node mapping:

Node-to-Edge mapping:



GAE Model for Graph Refinement: Encoder

Encoder: $\mathbf{h}_j^1 = g_n(\mathbf{x}_j)$ Node Embedding: $\mathbf{h}_j^1 = g_n(\mathbf{x}_j)$ Node-to-Edge mapping: $\mathbf{h}_{(i,j)}^1 = g_{n2e}([\mathbf{h}_i^1, \mathbf{h}_j^1])$ Edge-to-Node mapping: $\mathbf{h}_j^2 = g_{e2n}(\sum_{i}^{N_j} \mathbf{h}_{(i,j)}^1])$ Node-to-Edge mapping: $\mathbf{h}_{(i,j)}^2 = g_{n2e}([\mathbf{h}_i^2, \mathbf{h}_j^2])$

 $g_{...}(\cdot)$ are 2-layered MLPs, with ReLU, dropout and layer normalisation

GAE Model for Graph Refinement: Decoder

Decoder:

$$\alpha_{ij} = \sigma(g_{dec}(\mathbf{h}^2_{(i,j)})) \tag{2}$$

 g_{dec} is a 1D convolutional layer with one output channel Model can be trained by computing the loss $\mathcal{L}\alpha$, \mathbf{A}_r



Summarising GNNs



Outline

Airway Diseases

Output Description of the Data

Existing Methods Graph Refinement Model Mean-Field Networks Graph Neural Networks

4 Experiments



Training both models: Dice Loss

To tackle Class Imbalance:



Training both models: Dice Loss

To tackle Class Imbalance:

$$\mathcal{L}(\boldsymbol{\alpha}, \mathbf{A}_r) = 1 - \frac{2\sum_{i,j=1}^{N} \alpha_{ij} A_{ij}}{\sum_{i,j=1}^{N} \alpha_{ij}^2 + \sum_{i,j=1}^{N} A_{ij}^2},$$

 A_{ij} are individual binary entries \mathbf{A}_r



Data

- Danish Lung Cancer Screening Trial
- Low-dose Chest CT scans
- 32 scans with "Reference" annotations
- 100 scans with automatic segmentation



Results

- Error Measure based on centerline distances
- Average of two distances, $d_{err} = (d_{FP} + d_{FN})/2$
- Compared with Top Performer on Airway Extraction Challenge

Method	$d_{FP}(\text{mm})$	$d_{FN}(\mathrm{mm})$	d_{err} (mm)
Voxel Classifier MFN GNN	$3.871 \\ 3.716 \\ 3.513$	$5.108 \\ 3.992 \\ 2.890$	$\begin{array}{c} 4.489 \pm 0.754 \\ 3.845 \pm 0.415 \\ 3.202 \pm 0.386 \end{array}$

Table I: Performance comparison

