

Online Graph Topology Inference with Kernels

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Structure

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- 2 Problem definition and optimization
- 3 Experiment & Results
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Motivation

- data are abundant and diverse, and are often supported by irregular domains that can be naturally modeled as graphs
- most graph signal processing algorithms assume prior knowledge of the graph structure
- examples where topology needs to be inferred from data include brain networks, gene regulation systems [1] or social and economical interactions [2]



Figure 1: Potential brain network.
(Sapien Labs)

Definitions and notations



Definitions

- graph $\mathcal{G} = (\mathcal{N}, \mathcal{E})$
- \mathcal{N} set of $N + 1$ nodes
- \mathcal{E} set of edges; if m and n are linked, $(m, n) \in \mathcal{E}$
- *adjacency matrix* \mathbf{A} [3, 4]
 - $(N + 1) \times (N + 1)$ matrix
 - a_{nm} is 0 if $(m, n) \notin \mathcal{E}$, 1 otherwise
 - encodes the underlying graph connectivity
- signal $\mathbf{y}(i) \triangleq [y_1(i), \dots, y_{N+1}(i)]^T, i \in \mathbb{N}_+$

Background



Goal

Estimating the graph topology encoded in a (possibly directed) adjacency matrix \mathbf{A} from online nodal measurements $\mathbf{y}(i) = [y_1(i), \dots, y_{N+1}(i)]^T$, $i \in \mathbb{N}_+$ acquired over \mathcal{G}

- The dynamic graph signal $\mathbf{y}(i)$ can denote, e.g., the **electrical activity of different brain-regions** [5, 6], or the **voltage angle per bus** [7]
- The signal at each node $y_n(i)$ influences and is influenced by the signals at the other nodes ($y_m(i)$, $m \in \mathcal{N} \setminus \{n\}$)

Additive signal model

Nonlinear interactions are being reported in many applications (e.g., in brain connectivity [8, 9])

Recent methods have considered models of the form [10]:

$$y_n(i) = \sum_{m \in \mathcal{N} \setminus \{n\}} a_{nm} f_m(y_m(i)) + v_n(i), \quad (1)$$

where

- a_{nm} is the $(n, m)^{\text{th}}$ entry of the graph adjacency matrix \mathbf{A}
- f_m is a nonlinear function
- $v_n(i)$ represents innovation noise

Optimization problem in the input space

Using model (1), the topology estimation problem can be formulated using all available measurements $(y_n(\ell))$ for $\ell \leq i$ as:

$$\begin{aligned} \underset{\mathbf{a}_n, f_1, \dots, f_N}{\operatorname{argmin}} \quad & \frac{1}{2i} \sum_{\ell=1}^i \left\| y_n(\ell) - \sum_{m \in \mathcal{N} \setminus \{n\}} a_{nm} f_m(y_m(\ell)) \right\|^2 + \vartheta(\mathbf{a}_n) \\ \text{subject to} \quad & a_{nm} \in \{0, 1\}, \end{aligned} \quad (2)$$

where \mathbf{a}_n is the n^{th} row of \mathbf{A} , and function ϑ is a sparsity promoting regularization (e.g., ℓ_0 or ℓ_1 (semi)-norm).

However, problem (2) is difficult to solve: it is non-convex, and has infinite dimensional decision variables f_m .

Optimization problem in the RKHS

To obtain an efficient algorithm without sacrificing representation power, we:

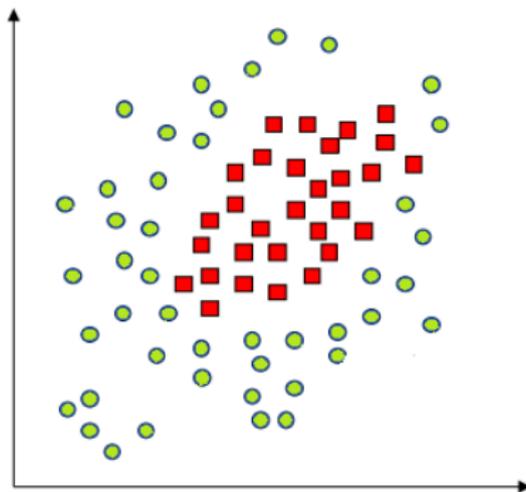
- denote $\phi_{nm} = a_{nm} f_m$, which allows us to incorporate the binary variable a_{nm} and turn (2) into a problem that is quadratic in ϕ_{nm} .
- constrain ϕ_{nm} , for $m \in \mathcal{N} \setminus \{n\}$, to a Reproducing Kernel Hilbert Space (RKHS) \mathcal{H}_κ associated with a positive definite reproducing kernel $\kappa(\cdot, \cdot)$.
- Thus, $a_{nm} = 0$ becomes equivalent to $\|\phi_{nm}\|_{\mathcal{H}_\kappa} = 0$.

The optimization problem becomes:

$$\underset{\substack{\phi_{nm} \in \mathcal{H}_\kappa \\ m=1, \dots, N}}{\operatorname{argmin}} \frac{1}{2i} \sum_{\ell=1}^i \left\| y_n(\ell) - \sum_{m \in \mathcal{N} \setminus \{n\}} \phi_{nm}(y_m(\ell)) \right\|^2 + \sum_{m \in \mathcal{N} \setminus \{n\}} \psi_{\mathcal{H}_\kappa}(\|\phi_{nm}\|_{\mathcal{H}_\kappa}), \quad (3)$$

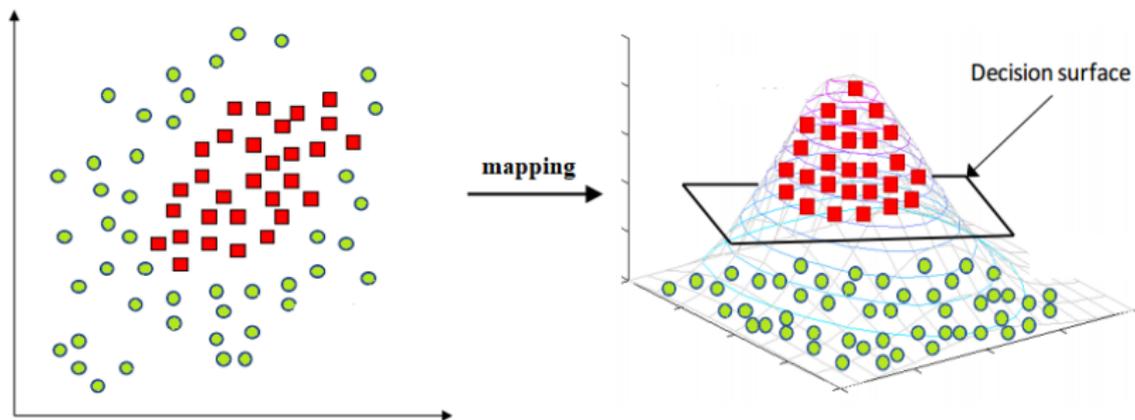
where $\psi_{\mathcal{H}_\kappa} : \mathbb{R} \rightarrow [0, \infty[$ is a non-decreasing function.

The Kernel Trick



<https://medium.com/@zxr.nju/what-is-the-kernel-trick-why-is-it-important-98a98db0961d>

The Kernel Trick



$$\begin{aligned} \text{Mapping } \varphi : \mathbb{R}^{d_1} &\rightarrow \mathbb{R}^{d_2}, d_2 \gg d_1, \mathbf{x} \in \mathbb{R}^{d_1} \mapsto \varphi(\mathbf{x}) \in \mathbb{R}^{d_2} \\ \implies \forall f(\cdot) \in \mathbb{R}^{d_2}, f(\mathbf{x}) &= \langle f(\cdot), \kappa(\cdot, \mathbf{x}) \rangle_{\mathbb{R}^{d_2}}, \text{ with } \kappa(\cdot, \mathbf{x}) = \varphi(\mathbf{x}) \\ &\implies \kappa(\mathbf{x}_1, \mathbf{x}_2) = \langle \varphi(\mathbf{x}_1), \varphi(\mathbf{x}_2) \rangle_{\mathbb{R}^{d_2}} \end{aligned}$$

$$\text{Examples: } \kappa(\mathbf{x}_1, \mathbf{x}_2) = \exp(-\gamma \|\mathbf{x}_1 - \mathbf{x}_2\|^2), \kappa(\mathbf{x}_1, \mathbf{x}_2) = \tanh(a\mathbf{x}_1^\top \mathbf{x}_2 + b)$$

<https://medium.com/@zxr.nju/what-is-the-kernel-trick-why-is-it-important-98a98db0961d>

Finite dimensional representation

The representer theorem [11] implies that the solution to (3) admits a finite-dimensional representation:

$$\phi_{nm}^*(\cdot) = \sum_{p=1}^i \alpha_{nmp} \kappa_m(\cdot, y_m(p)), \quad m = 1, \dots, N, \quad \alpha_{nmp} \in \mathbb{R} \quad (4)$$

However, the number of coefficients $\{\alpha_{nmp}\}$ increases with i , which is a problem for online processing.

Sparse kernel dictionaries

A solution to this problem is to consider sparse kernel dictionaries \mathcal{D}_m :

Kernel dictionary and sparsification rule

- each node m in the network creates, updates, and stores a dictionary of kernel functions, $\mathcal{D}_m = \{\kappa_m(\cdot, y_m(\omega_j)) : \omega_j \in \mathcal{I}_m^i \subset \{1, \dots, i-1\}\}$
- a candidate kernel function $\kappa_m(\cdot, y_m(i))$ is added in \mathcal{D}_m if the following sparsification condition holds [12]:

$$\max_{\omega_j \in \mathcal{I}_m^i} |\kappa_m(y_m(i), y_m(\omega_j))| \leq \xi_m, \quad (5)$$

where $\xi_m \in [0, 1[$ determines the level of sparsity and coherence [12]

- the size of the dictionary remains bounded as $i \rightarrow \infty$

Introducing sparsity

Since ϕ_{nm} encodes the interaction from node m to node n in model (1), promoting sparsity over $\mathbf{A} \Leftrightarrow$ promoting sparsity over the functions ϕ_{nm} .

The coefficient-based representation (4) means that sparsity of groups of variables $\{\alpha_{nm\omega_j}\}_{\omega_j \in \mathcal{I}_m^i}$, $m \in \mathcal{N} \setminus \{n\}$, can be promoted by using a block-sparse regularization [13]:

$$\alpha_n^* = \underset{\alpha_n}{\operatorname{argmin}} \frac{1}{2} \left\| y_n(i) - \alpha_n^\top \tilde{\mathbf{k}}(i) \right\|^2 + \eta_n \sum_{m \in \mathcal{N} \setminus \{n\}} \|\tilde{\alpha}_{nm}\|_2, \quad (6)$$

where we considered the instantaneous MSE estimate (measured only at instant i), with block vectors α_n and $\tilde{\mathbf{k}}(i)$ are defined as:

$$\begin{aligned} \tilde{\mathbf{k}}(i) &= [\mathbf{k}_1^\top(i), \dots, \mathbf{k}_N^\top(i)]^\top, \quad \mathbf{k}_m(i) = \operatorname{col}\{k_m(y_m(i), y_m(\omega_j))\}_{\omega_j \in \mathcal{I}_m^i}, \\ \alpha_n &= [\tilde{\alpha}_{n1}^\top, \dots, \tilde{\alpha}_{nN}^\top]^\top, \quad \tilde{\alpha}_{nm} = \operatorname{col}\{\alpha_{nm\omega_j}\}_{\omega_j \in \mathcal{I}_m^i}. \end{aligned} \quad (7)$$

Algorithm update

Using the subgradient descent algorithm [14] leads to the update:

Update rule

$$\hat{\alpha}_n(i+1) = \hat{\alpha}_n(i) + \mu_n \tilde{\mathbf{k}}(i) [y_n(i) - \tilde{\mathbf{k}}^\top(i) \hat{\alpha}_n(i)] - \mu_n \eta_n \mathbf{\Gamma}_n(i). \quad (8)$$

with $\mathbf{\Gamma}_n(i) = [\mathbf{\Gamma}_{n1}^\top(i), \dots, \mathbf{\Gamma}_{nN}^\top(i)]^\top$ [14], where each block $\mathbf{\Gamma}_{nm}(i)$ is:

$$\mathbf{\Gamma}_{nm}(i) = \begin{cases} \frac{\tilde{\alpha}_{nm}(i)}{\|\tilde{\alpha}_{mn}(i)\|_2} & \text{if } \|\tilde{\alpha}_{mn}(i)\|_2 \neq 0 \\ \mathbf{0} & \text{if } \|\tilde{\alpha}_{mn}(i)\|_2 = 0 \end{cases}. \quad (9)$$

Edge identification

Set $\hat{a}_{nm}(i)$ to 1 if $\|\hat{\alpha}_{nm}(i)\| \geq \tau_n$, to 0 otherwise

Epilepsy dataset setting

The used data come from a 39-year-old female subject suffering from intractable epilepsy [15]. The data-set contains 8 instances of electrocorticography (ECoG) time series, each instance representing one seizure and contains voltage measurements from 76 different regions on and inside the brain, during:

- the 10 seconds before the epilepsy seizure (*preictal* interval)
- the first 10 seconds during the seizure (*ictal* interval)

Epilepsy dataset results

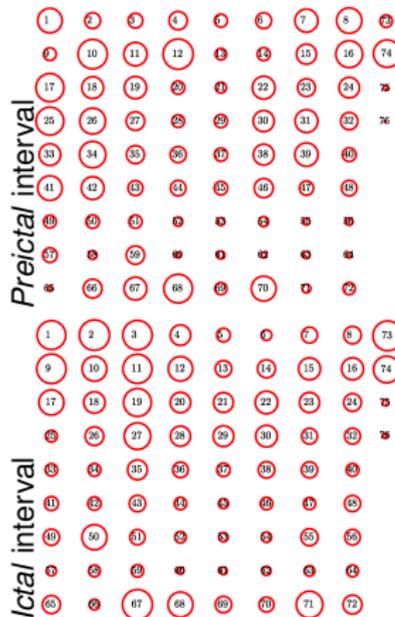
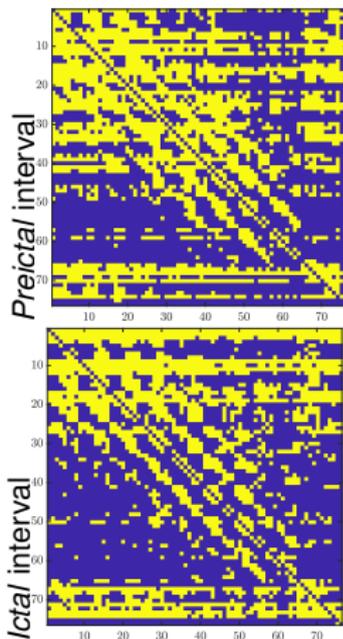


Figure 2: Estimated adjacency matrices (left) and summed in- and out-degrees for the estimated graphs (right). The larger the radius corresponding to node n , the larger the summed degree of node n .

Conclusion and perspectives

Conclusion

- online adaptive graph topology algorithm
- the use of kernels allows for inferring nonlinear relationships
- kernel dictionaries mitigate the increasing number of data points inherently present in an online setting
- consistent results on real data

Perspectives

- the use of multi-kernels
- the use of other sparsity-inducing techniques

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Schizophrenia dataset setting



We used electroencephalography (EEG) measurements [16] taken from a group of six subjects, half of which are healthy and half suffer from schizophrenia. A simple button-pressing task is set up, in three separate settings where subjects either:

- 1 Task 1: pressed the button and a tone was immediately played
- 2 Task 2: listened to the tone without the button press
- 3 Task 3: pressed the button and the tone was not played

Schizophrenia dataset results

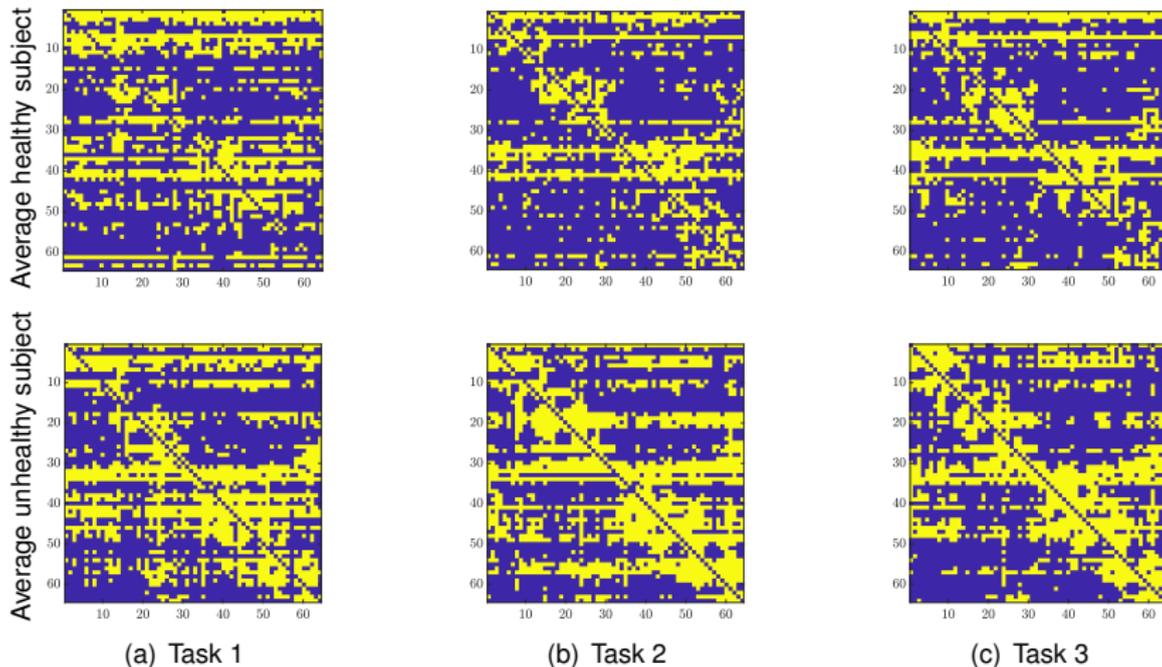


Figure 3: Estimated topologies per task, averaged per group.