

# Directed networks via Persistent Hochschild Homology

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# Background

Network data in many guises in life sciences:

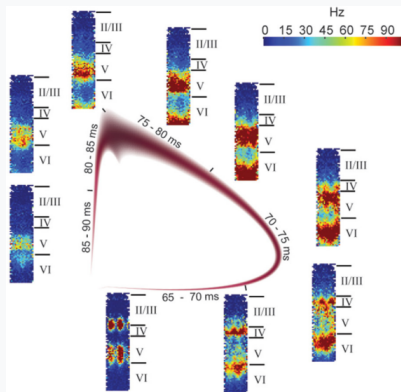
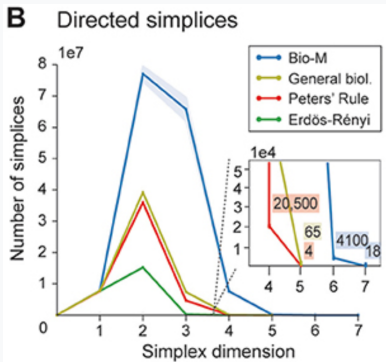
- protein–protein interaction networks
- gene regulatory networks
- vascular network
- neuronal networks

From TDA perspective, what is an appropriate (persistent) homology theory for networks, or directed graphs?

- use homological information to understand some aspects of, for example, network dynamics

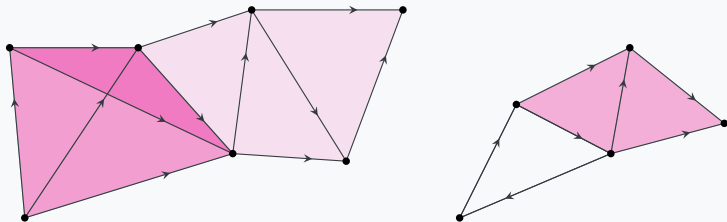
# Some topological observations in neuronal networks

(M. Reimann et al., *Cliques of Neurons Bound into Cavities Provide a Missing Link between Structure and Function*, 2017)

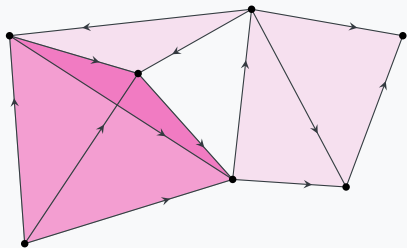
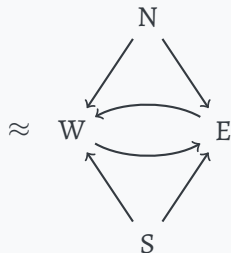
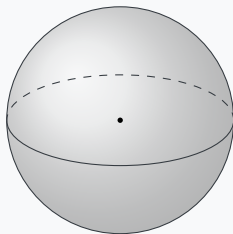
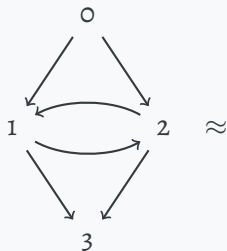


# Directed flag complex of a digraph $G$

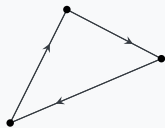
$n$ -simplices are totally ordered sets of vertices  $v_0 < v_1 < \dots < v_n$ , such that for any  $v_i < v_j$ , the pair  $(v_i, v_j)$  is a directed edge in  $G$ .



# Topology of directed flag complexes



$\approx$

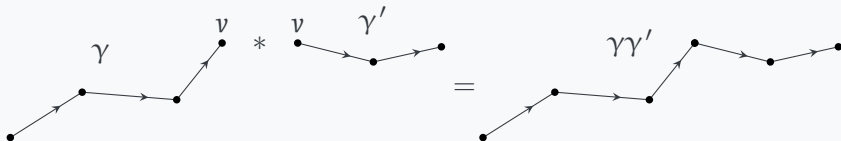


# Homology from directed paths?

From network analysis perspective, phenomena travel over the directed edges and paths. *We want homological tools to account for this.*

The **path algebra**  $kG$  over a field  $k$  associated to the digraph  $G$

- has as a  $k$ -vector space a basis consisting of all possible paths  $\gamma$  in  $G$
- and the multiplication is concatenation of paths end-to-end if possible, and 0 otherwise.



## Path algebra and Hochschild homology

We can use **Hochschild homology** of associative algebras as a homology theory for directed graphs.

$$\dots \xrightarrow{b_{n+1}} kG^{\otimes n+1} \xrightarrow{b_n} kG^{\otimes n} \xrightarrow{b_{n-1}} \dots \xrightarrow{b_1} kG.$$

Happel's combinatorial formula for Betti numbers when  $G$  is a connected, acyclic, directed graph:

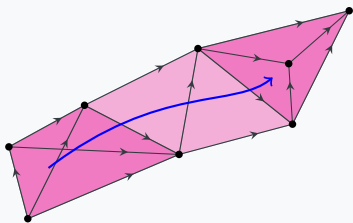
$$\dim_k HH_i(kG) = \begin{cases} 1 & \text{if } i = 0 \\ 0 & \text{if } i > 1 \\ 1 - n + \sum_{e \in G} \dim_k s_e kGt_e & \text{if } i = 1, \end{cases}$$

where  $n$  is the number of vertices of  $G$  and  $s_e kGt_e$  is the subspace generated by all the paths from start of  $e$  to end of  $e$ . *No higher degree homological information.*

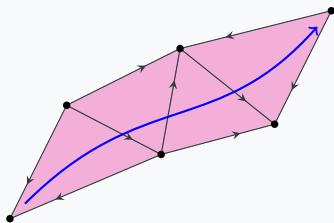
# Higher Hochschild homology of digraphs

We introduce higher degree  $HH$  by lifting to paths of simplices.

A **connectivity structure** of a digraph  $E: \text{Sub}(G) \times \text{Sub}(G) \rightarrow \{0, 1\}$ .



for simplices  $\sigma$  and  $\tau$   
and directions  $(d_i, d_j)$ ,  
 $(\sigma, \tau) \mapsto 1$  if either  $\sigma \hookrightarrow \tau$   
or  $d_i(\sigma) \hookrightarrow \alpha \hookrightarrow d_j(\tau)$ ,  
for some  $q$ -simplex  $\alpha$



for  $n$ -simplices  $\sigma$  and  $\tau$ ,  
 $(\sigma, \tau) \mapsto 1$  if there is  
an  $(n-1)$ -simplex  $\alpha$  and some  
 $i, j \in \{0, \dots, n\}$  such that  
 $d_i(\sigma) = \alpha = d_j(\tau)$ , with  $i < j$



# Persistent Hochschild homology from connectivity digraphs

1. For a digraph  $G$ , apply connectivity structure  $E$  to get a map

$$\mathbf{Digraph} \xrightarrow{E} \mathbf{Digraph}(\text{of simplices})$$

2. Go to the path algebra

$$\mathbf{Digraph} \xrightarrow{E} \mathbf{Digraph}(\text{of simplices}) \xrightarrow{k-} k\text{-Alg}$$

3. Compute Hochschild homology of the path algebra

$$\mathbf{Digraph} \xrightarrow{E} \mathbf{Digraph}(\text{of simplices}) \xrightarrow{k-} k\text{-Alg} \xrightarrow{HH} \mathbf{FinVect}$$

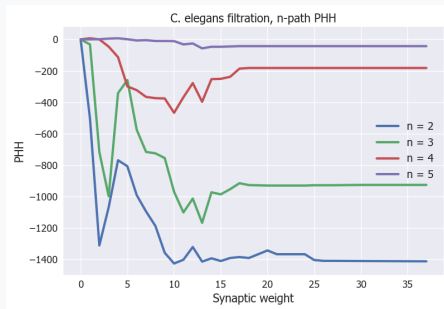
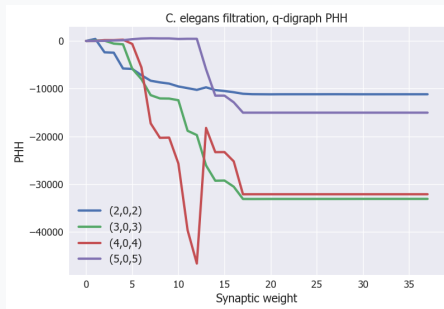
4. Adding filtration and (non-functorial) condensation

$$(\mathbf{R}, \leq) \xrightarrow{\mathcal{F}} \mathbf{Digraph} \xrightarrow{E} \mathbf{Digraph} \xrightarrow{c} \mathbf{Digraph}_{\text{acyc}} \xrightarrow{k-} k\text{-Alg} \xrightarrow{HH} \mathbf{FinVect}$$

When the pipeline is fully functorial, we have a stability theorem.

# Persistent *HH* of *C. elegans* neuronal network

Filtration induced by the synaptic strength between neurons. The erratic behaviour is due to the condensation operation.



Fin.

# Thanks!

Henri Riihimäki, *Simplicial  $q$ -connectivity of directed graphs with applications to network analysis*, to appear in SIAM Journal on Mathematics of Data Science, arXiv:2202.07307, 2022.

Luigi Caputi, Henri Riihimäki, *Hochschild homology, and a persistent approach via connectivity digraphs*, Journal of Applied and Computational Topology, 2023.

L. Caputi and H. Riihimäki, *On reachability categories and commuting algebras of quivers*, arXiv:2306.15388, 2023.