







# Learning with Covariance Matrices: Foundations and Applications to Network Neuroscience

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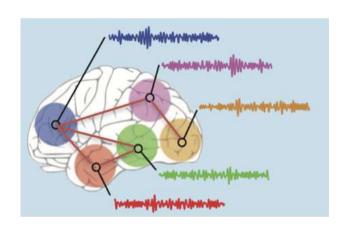
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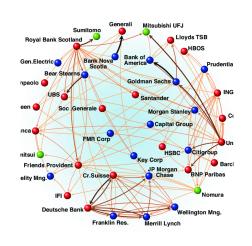
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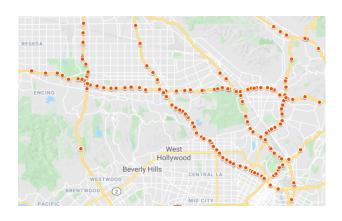


#### Covariance Matrix

- > Covariance matrix captures the **redundancies** between data points (features)
  - Brain datasets: some areas of the brain activate together
  - Financial datasets: stock prices fluctuate in tandem
  - Traffic datasets: traffic volume is correlated across intersections





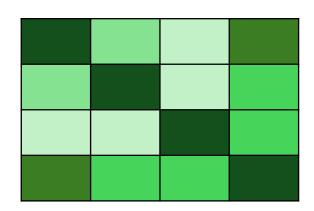


Finance Traffic

Brain

### Covariance Matrix

- > Evaluating a covariance matrix
  - Consider a random variable  $\mathbf{x} \in \mathbb{R}^m$
  - The covariance is



$$\mathbf{C} = \mathbb{E}[(\mathbf{x} - \boldsymbol{\mu})(\mathbf{x} - \boldsymbol{\mu})^{\mathsf{T}}], \text{ where } \boldsymbol{\mu} = \mathbb{E}[\mathbf{x}]$$

In practice, we have sample covariance matrix (an estimate)

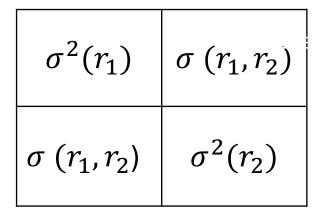
$$\hat{\mathbf{C}} = \frac{1}{n-1} \sum_{i=1}^{n} (\mathbf{x}_i - \hat{\boldsymbol{\mu}}) (\mathbf{x}_i - \hat{\boldsymbol{\mu}})^\mathsf{T}, \text{ where } \hat{\boldsymbol{\mu}} = \frac{1}{n} \sum_{i=1}^{n} \mathbf{x}_i$$

n: number of samples (size of a dataset)

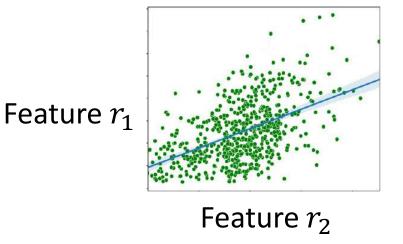
### Covariance Matrix

> Covariance matrix encodes redundancies between different features in data

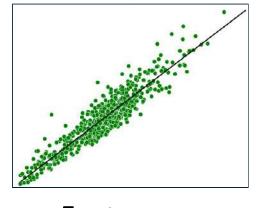
Covariance matrix (2-feature dataset)



**Low** redundancy (smaller  $\sigma$  ( $r_1$ ,  $r_2$ ))



**High** redundancy (higher  $\sigma(r_1, r_2)$ )



Feature  $r_1$ 

Feature  $r_2$ 

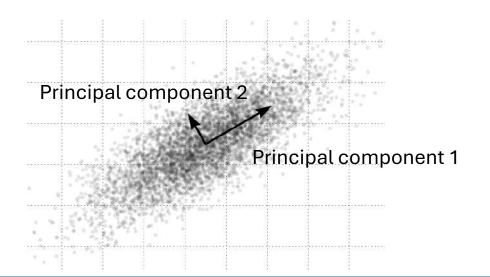
 $\sigma\left(r_{1},r_{2}\right)$  = how features  $r_{1}$  and  $r_{2}$  vary with respect to each other

#### Covariance matrices are widespread in signal processing and machine learning

- > Principal component analysis (PCA)
  - Eigenvectors of the covariance matrix form principal components (PCs)
  - PCs inform the shape of a dataset (directions of variance)

Given sample  ${\bf x}$  and eigendecomposition  $\hat{{\bf C}} = \hat{{\bf V}} \hat{{\bf \Lambda}} \hat{{\bf V}}^{\mathsf{T}}$ ,

PCA transform: 
$$\tilde{\mathbf{x}} = \hat{\mathbf{V}}^\mathsf{T} \mathbf{x}$$



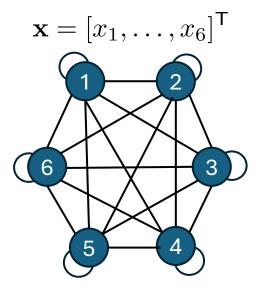
#### PCA transform in ML

- Unsupervised learning (dim. reduction)
- Supervised learning (regression, classification)

#### Covariance matrices are widespread in signal processing and machine learning

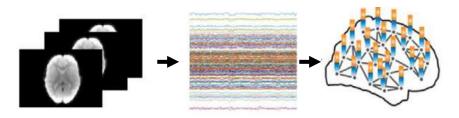
- > Covariance matrices are leveraged as graphical representations of data
  - A graph G = (V, E, W)
    - $_{\circ}$  Set of nodes V  $_{\circ}$  A weight function W
    - Set of edges *E*

- Covariance matrix is a fully connected graph,
  - nodes are the features
  - edges associated with pairwise covariance values

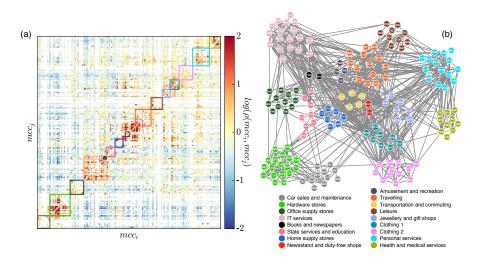


#### Covariance matrices are widespread in signal processing and machine learning

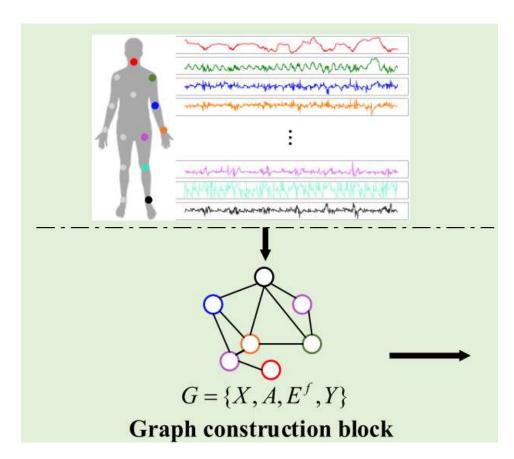
> Covariance matrices as graphical representations; used in graph neural nets



Brain connectome [Li, et al. 2021]



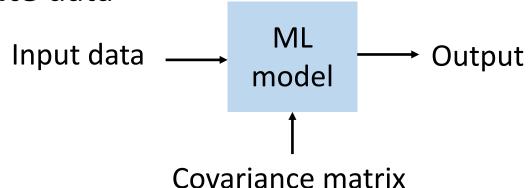
Socio-economic networks [Leo, et al. 2016]



Wearable devices [Wang, et al. 2023]

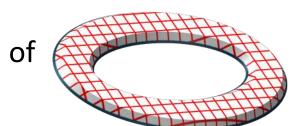
> Sample covariance matrix is estimate from **finite** data

- ML model is trained on training dataset, deployed on test dataset
- Statistical spaces defined by training and test data may not align perfectly



$$\hat{\mathbf{C}} = \frac{1}{n-1} \sum_{i=1}^{n} (\mathbf{x}_i - \hat{\boldsymbol{\mu}}) (\mathbf{x}_i - \hat{\boldsymbol{\mu}})^\mathsf{T}$$

Representation of training dataset



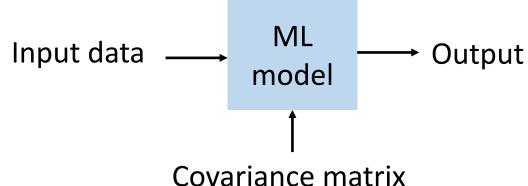
Representation of test dataset

> Sample covariance matrix is estimate from **finite** data

- > ML model is trained on training dataset, deployed on test dataset
- Statistical spaces defined by training and test data may not align perfectly

#### **Challenge 1 (stability)**

Are inference outcomes **stable** to perturbations in covariance matrix (finite sample effect)?

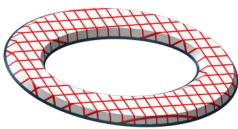


 $\hat{\mathbf{C}} = \frac{1}{n-1} \sum_{i=1}^{n} (\mathbf{x}_i - \hat{\boldsymbol{\mu}}) (\mathbf{x}_i - \hat{\boldsymbol{\mu}})^\mathsf{T}$ 

Representation of training dataset

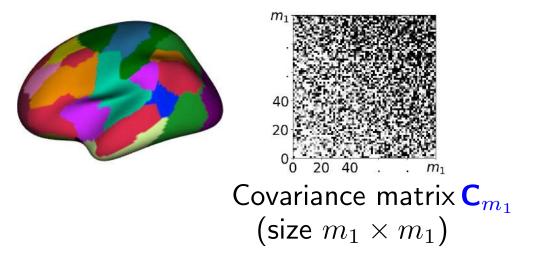


Representation of test dataset

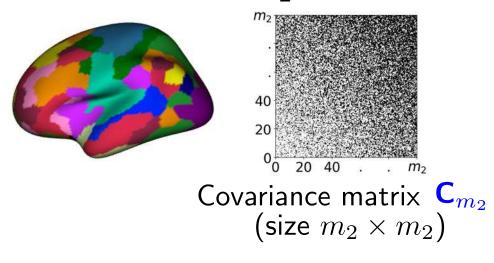


Datasets capture information about same phenomenon at different scales

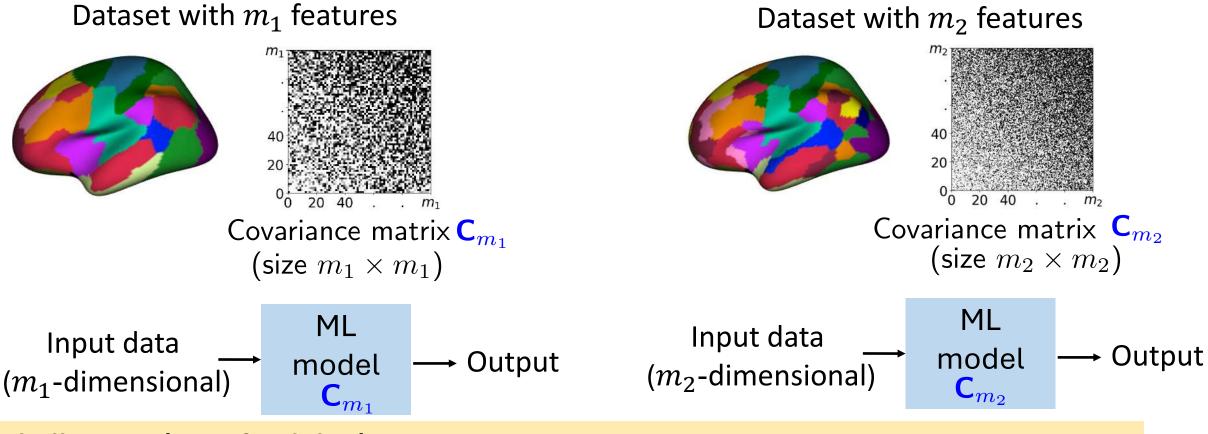
Dataset with  $m_1$  features



Dataset with  $m_2$  features



Datasets capture information about same phenomenon at different scales



#### **Challenge 2 (transferability)**

Can the redundancy in covariance matrices of datasets of different sizes be exploited?

# Learning with covariance matrices: A GSP approach

- > Signal and information processing is about exploiting signal structure
- > Graph signal processing (GSP): broaden classical signal processing to graphs



#### **Graph Signal Processing:** Overview, Challenges, and **Applications**

This article presents methods to process data associated to graphs (graph signals) extending techniques (transforms, sampling, and others) that are used for conventional signals.

By Antonio Ortega<sup>®</sup>, Fellow IEEE, Pascal Frossard, Fellow IEEE, Jelena Kovačević, Fellow IEEE. IOSÉ M. F. MOURA . Fellow IEEE, AND PIERRE VANDERGHEYNST

Graphs offer the ability to model such data and complex

interactions among them. For example, users on Twitter can be

modeled as nodes while their friend connections can be modeled

as edges. This paper explores adding attributes to such nodes and

modeling those as signals on a graph; for example, year of gradua

tion in a social network, temperature in a given city on a given day

in a weather network, etc. Doing so requires us to extend classical

signal processing concepts and tools such as Fourier transform,

filtering, and frequency response to data residing on graphs. It

also leads us to tackle complex tasks such as sampling in a princi-

pled way. The field that gathers all these questions under a com-

given later in the paper, let us assume for now that a graph

signal is a set of values residing on a set of nodes. These nodes

are connected via (possibly weighted) edges. As in classical

signal processing, such signals can stem from a variety of

domains; unlike in classical signal processing, however, the

underlying graphs can tell a fair amount about those signals

through their structure. Different types of graphs model dif-

world data include Erdős-Rényi graphs, ring graphs, random

geometric graphs, small-world graphs, power-law graphs,

nearest-neighbor graphs, scale-free graphs, and many others.

These model networks with random connections (Erdős-

graphs), social networks (scale-free graphs), and others.

Rényi graphs), networks of brain neurons (small-world

properties, such as smoothness, that need to be appropri-

ately defined. They can also be represented via basic atoms

and can have a spectral representation. In particular, the

graph Fourier transform allows us to develop the intuition

gathered in the classical setting and extend it to graphs; we

can talk about the notions of frequency and bandlimitedness

As in classical signal processing, graph signals can have

Typical graphs that are used to represent common real-

ferent types of networks that these nodes represent.

While the precise definition of a graph signal will be

mon umbrella is graph signal processing (GSP) [2], [3].

ABSTRACT | Research in graph signal processing (GSP) aims to develop tools for processing data defined on irregular graph domains. In this paper, we first provide an overview of core ideas in GSP and their connection to conventional digital signal processing, along with a brief historical perspective to highlight how concepts recently developed in GSP build on top of prior research in other areas. We then summarize recent advances in developing basic GSP Next, we review progress in several application areas using GSP, including processing and analysis of sensor network data, biological data, and applications to image processing and machine learning.

KEYWORDS | Graph signal processing (GSP); network science and graphs; sampling; signal processing

#### I. INTRODUCTION AND MOTIVATION

Data is all around us, and massive amounts of it. Almost every aspect of human life is now being recorded at all levels: from the marking and recording of processing inside the cells starting with the advent of fluorescent markers, to our personal data through health monitoring devices and apps, financial and banking data, our social networks, mobility and traffic patterns, marketing preferences, fads, and many more. The complexity of such networks [1] and interactions. means that the data now reside on irregular and complex structures that do not lend themselves to standard tools.

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P. Frossard, and P. Vandergheynst are with EPFL, Lausanne, Switzerland-1015,

0018-9219 © 2018 IEEE, Personal use Authorized licensed use limited to "INIVERSITY AT ALEANY, Downloader on August 22 2025 at 19 32 TO UTC from TEEE Xplore. Restrictions apply 808 PROCEEDINGS OF THE IEEE I Vol. 106, No. 5, May 2018 Geert Leus<sup>®</sup>, Antonio G. Marques<sup>®</sup>, José M.F. Moura<sup>®</sup>, Antonio Ortega<sup>®</sup>, and David I Shuman<sup>®</sup>

75TH ANNIVERSARY OF SIGNAL PROCESSII

#### **Graph Signal Processing**

History, development, impact, and outlook



nferring information defined over regular (first continu ous, later discrete) domains such as time or space. Indeed he last 75 years have shown how SP has made an impact in areas such as communications, acoustics, sensing, image processing, and control, to name a few. With the digitalization of the modern world and the increasing pervasiveness of data-collection mechanisms, information of interest in current nains. Graph SP (GSP) generalizes SP tasks to signals living on non-Euclidean domains whose structure can be captured by a weighted graph. Graphs are versatile, able to model irreguir interactions, easy to interpret, and endowed with a corpus of mathematical results, rendering them natural candidates to serve as the basis for a theory of processing signals in more rregular domains The term graph signal processing was coined a decade ago

n the seminal works of [1], [2], [3], and [4]. Since these papers were published, GSP-related problems have drawn significant attention, not only within the SP community [5] but also in machine learning (ML) venues, where research in graph-based learning has increased significantly [6]. Graph signals are wellsuited to model measurements/information/data associated with (indexed by) a set where 1) the elements of the set belon to the same class (regions of the cerebral cortex, members of a social network, weather stations across a continent): 2) there exists a relation (physical or functional) of proximity, influence or association among the different elements of that set; and 3) not homogeneous. In some scenarios, the supporting graph is a physical, technological, social, information, or biological net work where the links can be explicitly observed. In many other cases, the graph is implicit, capturing some notion of dependence or similarity across nodes, and the links must be inferred from the data themselves. As a result, GSP is a broad frame work that encompasses and extends classical SP methods, tools and algorithms to application domains of the modern technological world, including social, transportation, communication,

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Xiaowen Dong, Dorina Thanou, Laura Toni,

#### **Graph Signal Processing for Machine Learning**

A review and new perspectives



ne effective representation, processing, analysis, and visual ization of large-scale structured data, especially those related to complex domains, such as networks and graphs, are one of the key questions in modern machine learning. Graph signal processing (GSP), a vibrant branch of signal processing models and algorithms that aims at handling data supported on graphs, onens new paths of research to address this challenge. In this article, we review a few important contributions made by GSP concepts and tools, such as graph filters and transforms, to the devel opment of novel machine learning algorithms. In particular, ou discussion focuses on the following three aspects: exploiting data structure and relational priors, improving data and computational efficiency, and enhancing model interpretability. Furthermore we provide new perspectives on the future development of GSP echniques that may serve as a bridge between applied mathematics and signal processing on one side and machine learning and network science on the other. Cross-fertilization across thes different disciplines may help unlock the numerous challenges of complex data analysis in the modern age.

We live in a connected society. Data collected from large-scale interactive systems, such as biological, social, and financial networks, become largely available. In parallel, the past few decades have seen a significant amount of interest in the machine learning community for network data processing and analysis. Networks have an intrinsic structure that conveys very specific properties to data, e.g., interdependencies between data entities in the form of pairwise relationships. These properties are traditionally captured by mathematical representations such as graphs.

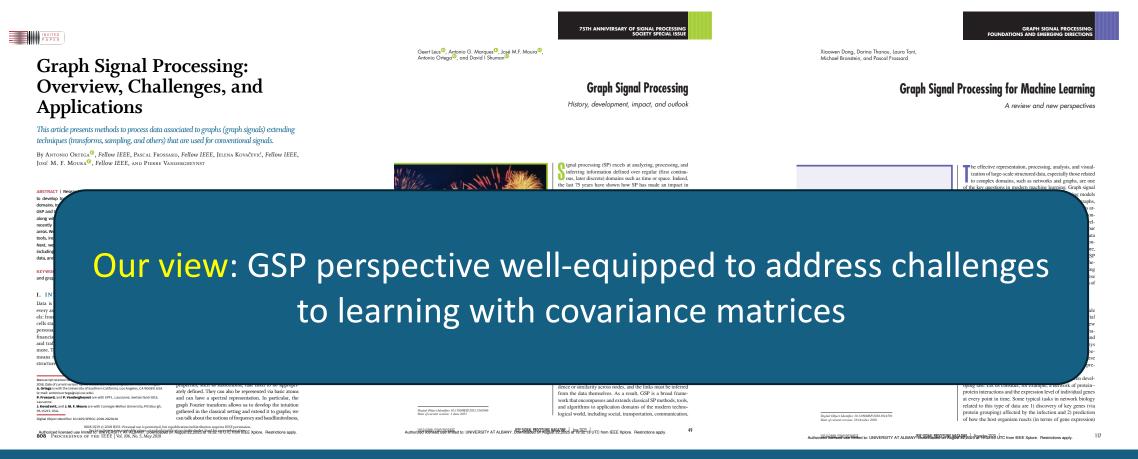
In this context, new trends and challenges have been developing fast. Let us consider, for example, a network of proteinprotein interactions and the expression level of individual genes at every point in time. Some typical tasks in network biology related to this type of data are 1) discovery of key genes (via protein grouping) affected by the infection and 2) prediction of how the host organism reacts (in terms of gene expression)

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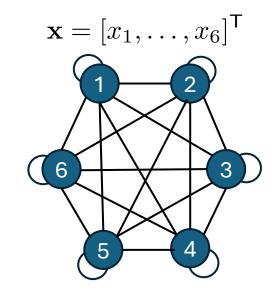
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- > Graph signal processing (GSP): broaden classical signal processing to graphs



# Learning with covariance matrices: A GSP approach

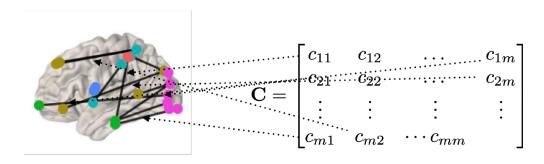
- > Graph neural networks (GNNs) have been shown to be [Ruiz et al., 2023]
  - stable to (abstract) perturbations in graph structure
  - generalizable to graph structures of different sizes (similar to convolutional neural nets for images)



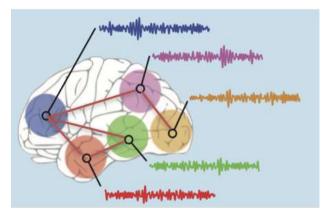
- Covariance matrix is a data-driven graph
  - interplay between perturbation theory of covariances and ML over them

## Applications to network neuroscience

> Covariance matrices appear commonly in network neuroscience



Anatomical covariance matrix

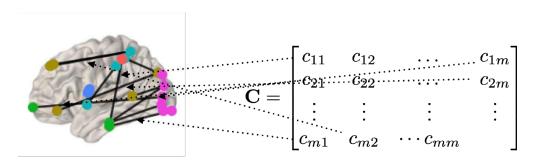


Functional connectome

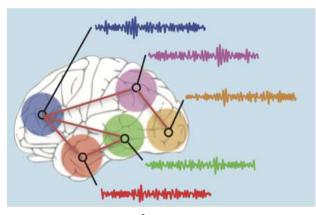
> Principled ML approaches for reproducible, transparent, generalizable findings

# Applications to network neuroscience

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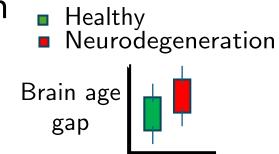


Anatomical covariance matrix



Functional connectome

- > Principled ML approaches for reproducible, transparent, generalizable findings
- > Brain age gap is a biomarker that reflects neurodegeneration
  - How VNN theoretical advances provide principled
     brain age gap prediction?



#### coVariance neural networks

> coVariance neural networks (VNNs):

GNNs operating on covariance matrices



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Disentangling Neurodegeneration With Brain Age Gap Prediction Models

A graph signal processing perspective

- Two tutorial articles in IEEE SPM
  - Tutorial article on 'Disentangling neurodegeneration with brain age gap prediction models' (to appear in 2025)
  - Tutorial article on 'CoVariance Neural Networks:
     Principal Component Analysis Meets Learning
     with Graphs' (under preparation)



eurodegeneration is the progressive loss of structure or function of neurons in the brain. Reduction in cortical thickness or volume over time has been a workhorse metric used to assess neurodegeneration in clinical settings; see case study 1 in "Case Study 1: Cortical Atrophy Characterizes Neurodegeneration in Alzheimer's Disease" for a demonstration of cortical atrophy assessment in the context of Alzheimer's disease (AD) relative to healthy individuals [healthy cohort (HC) group]. Naturally, visual inspection of T1-weighted brain magnetic resonance imaging (MRI) images and associated MRI quantification products are used along with other biological measurements to make a "subjective" assessment about the brain health of an individual. These assessments tend to be subjective because they lack a deterministic relationship between an individual's health status and the absolute values of the metrics observed within MRI scans [1]. Moreover, such methods cannot adequately account for the statistical complexities inherent within neuroimaging datasets that capture neurodegeneration. In particular, neurodegeneration is a characteristic of the healthy aging process and various neurological disorders [2], exhibiting correlated patterns across brain regions. Such statistical factors motivate well the use of data-driven methods to characterize

Automating or improving the analyses of brain MRI images is appealing for several reasons; MRI is a noninvasive proce-

### Outline

- > PCA and the graph Fourier transform
- > CoVariance neural networks (VNNs)
- > Theory of VNNs: Stability and transferability
- > Application: Principled brain age gap prediction with VNNs
- > Variants of VNNs

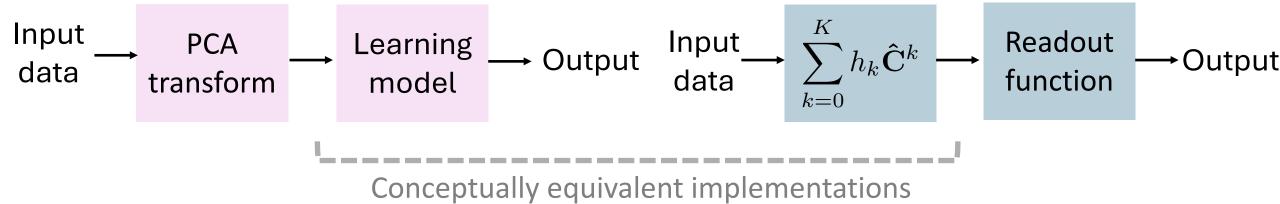
# Key takeaways

- VNNs offer a novel GSP-inspired perspective to PCA addressing challenges in modern data analysis
- Principled deep learning solution for finite-data regimes
  - Stability and transferability
- VNNs address methodological/conceptual obscurities in brain age gap prediction

### PCA and Graph Fourier Transform

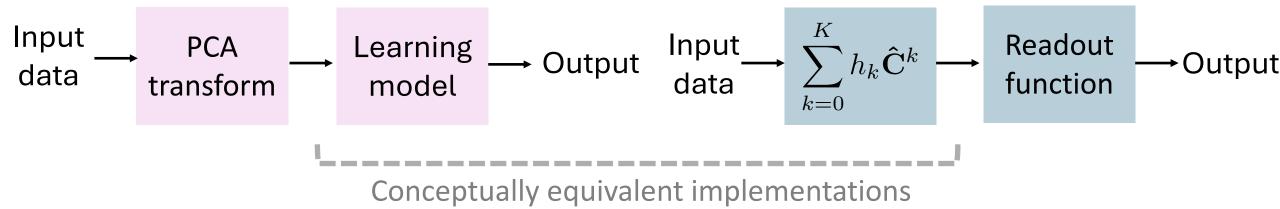
# A graph filter implementation of PCA inference

 $\succ$  To show: PCA-based inference can be implemented with a polynomial over  $\hat{\mathbf{C}}$ 



# A graph filter implementation of PCA inference

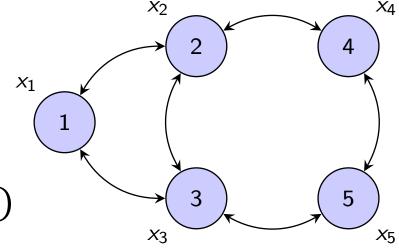
 $\succ$  To show: PCA-based inference can be implemented with a polynomial over  $\hat{\mathbf{C}}$ 



- **How:** Follows from the graph Fourier transform analysis of  $\sum_{k=0}^{\infty} h_k \hat{\mathbf{C}}^k$
- > Implications:
  - Alternative implementation of PCA-based inference using polynomial over  $\hat{\mathbf{C}}$
  - But more importantly, polynomial implementation is stable, transferable

# Preliminaries: Graph

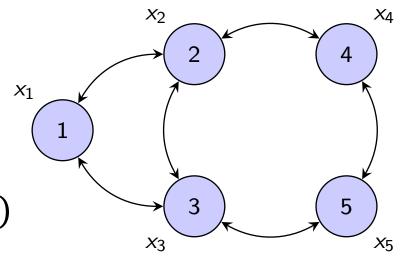
- $\triangleright$  **Graph:** a triplet (V, E, W)
  - A set of **nodes**  $V = \{1, ..., m\}$
  - A set of (undirected) **edges**  $E \subseteq V \times V$ Edge between node i and j denoted by (i,j)



• An **edge function**  $W: E \mapsto \mathbb{R}$  that maps edge (i, j)to weight  $w_{ij} \in \mathbb{R}$ 

# Preliminaries: Graph

- $\succ$  **Graph:** a triplet (V, E, W)
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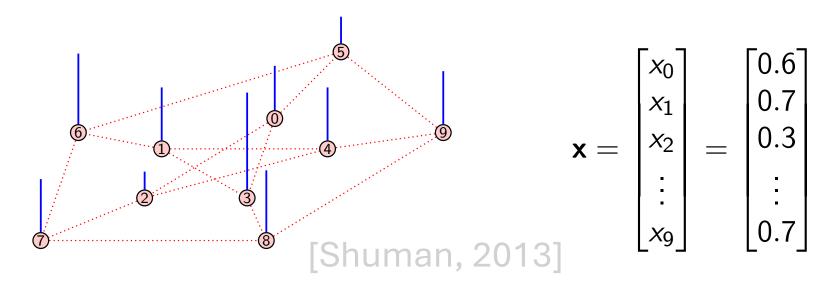


- An edge function  $W: E \mapsto \mathbb{R}$  that maps edge (i, j) to weight  $w_{ij} \in \mathbb{R}$
- Adjacency matrix representation of graph

$$[\mathbf{A}]_{ij} = \begin{cases} w_{ij}, & \text{if } (i,j) \in E, \\ 0, & \text{otherwise} \end{cases}$$

# Preliminaries: Graph signal

- $\triangleright$  Graph signals are mappings  $x:V\mapsto \mathbb{R}$ 
  - => graph signal is defined on the vertices of the graph
- $\succ$  **Graph signal** can be represented as a vector  $\mathbf{x} \in \mathbb{R}^m$ 
  - $\implies x_i$  denotes the graph signal at *i*-th vertex in V



# Preliminaries: Graph shift operator (GSO)

- $\succ$  To understand and analyze graph signal x, GSP accounts for the graph structure
- $\succ$  Graph structure is encoded in a graph shift operator  $\mathbf{S} \in \mathbb{R}^{m \times m}$

$$[S]_{ij} = 0$$
 for  $i \neq j$  and  $(i,j) \notin E$  (S captures local graph structure)

$$\mathbf{S} = \begin{pmatrix} S_{11} & S_{12} & 0 & 0 & S_{15} & 0 \\ S_{21} & S_{22} & S_{23} & 0 & S_{25} & 0 \\ 0 & S_{23} & S_{33} & S_{34} & 0 & 0 \\ 0 & 0 & S_{43} & S_{44} & S_{45} & S_{46} \\ S_{51} & S_{52} & 0 & S_{54} & S_{55} & 0 \\ 0 & 0 & 0 & S_{64} & 0 & S_{66} \end{pmatrix}$$

Examples: adjacency matrix, Laplacian

Covariance matrix is a data-driven adjacency matrix

# Preliminaries: Graph Fourier Transform (GFT)

- $\succ$  Generically, eigendecomposition of GSO  $S = U\Phi U^{-1}$
- > GFT is the projection of graph signal on the eigenvector space U

$$\tilde{\mathbf{x}} = \mathbf{U}^{-1}\mathbf{x}$$

Inverse GFT is defined as

$$\mathbf{x} = \mathbf{U} \, \tilde{\mathbf{x}}$$

 $\Longrightarrow$  Eigenvectors  $\mathbf{U} = [oldsymbol{u}_1, ..., oldsymbol{u}_m]$  are the frequency basis

### When GSO is covariance matrix...

> GFT over covariance matrix

Given eigendecomposition

$$\hat{\mathbf{C}} = \hat{\mathbf{V}} \hat{\mathbf{\Lambda}} \hat{\mathbf{V}}^\mathsf{T}$$

GFT of x is

$$\mathbf{\tilde{x}} = \hat{\mathbf{V}}^\mathsf{T} \mathbf{x}$$

### When GSO is covariance matrix...

> GFT over covariance matrix

Given eigendecomposition

$$\hat{\mathbf{C}} = \hat{\mathbf{V}} \hat{\mathbf{\Lambda}} \hat{\mathbf{V}}^\mathsf{T}$$

GFT of x is

$$\tilde{\mathbf{x}} = \hat{\mathbf{V}}^\mathsf{T} \mathbf{x}$$

> PCA transform

Projection of sample x on principal components of  $\hat{C}$ 

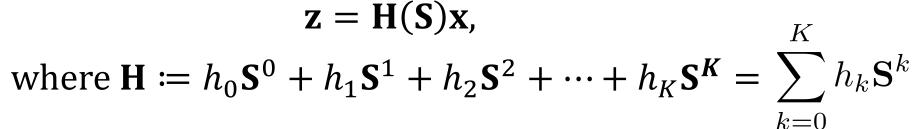
PCA transform: 
$$\tilde{\mathbf{x}} = \hat{\mathbf{V}}^\mathsf{T} \mathbf{x}$$

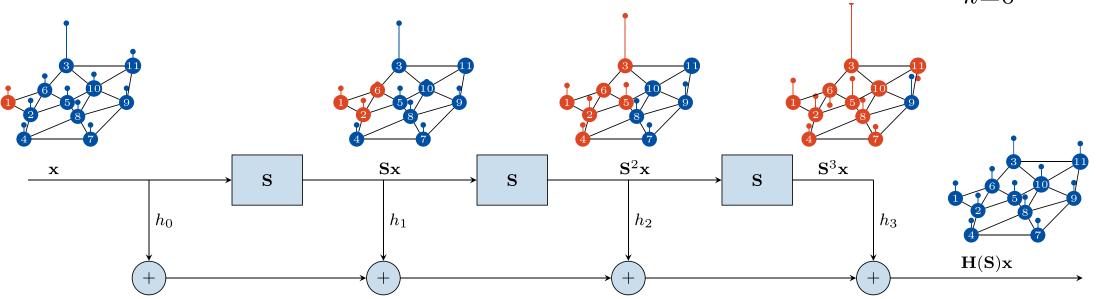
PCA transform is GFT with respect to the covariance graph!



# Preliminaries: Graph filter

ightharpoonup Graph filter H maps graph signal x to another graph signal z via linear-shift-and-sum operation



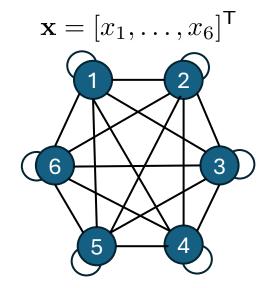


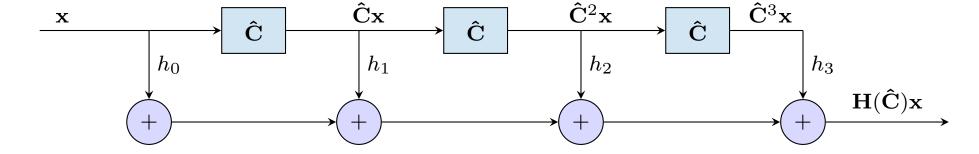
[Isufi et. al, IEEE TSP, 2024]

# Graph filter on covariance matrix

- > Covariance matrix forms a fully-connected graph where
  - nodes are features
  - edges are covariance values
- ightharpoonup Graph filter on covariance matrix  $\hat{\mathbf{C}}$  is defined as

$$\mathbf{H}(\hat{\mathbf{C}}) = \sum_{k=0}^{K} h_k \hat{\mathbf{C}}^k \mathbf{x}$$





#### CoVariance filter

- ightharpoonup Analogy between  $\mathbf{H}(\hat{\mathbf{C}})$  and PCA
  - Using eigendecomposition  $\,\hat{\mathbf{C}} = \hat{\mathbf{V}} \hat{\mathbf{\Lambda}} \hat{\mathbf{V}}^\mathsf{T}\,$  , it follows that

$$\mathbf{z} = \mathbf{H}(\hat{\mathbf{C}})\mathbf{x} = \sum_{k=0}^{K} h_k \hat{\mathbf{C}}^k \mathbf{x} = \sum_{k=0}^{K} h_k \hat{\mathbf{V}} \hat{\boldsymbol{\Lambda}}^k \hat{\mathbf{V}}^\mathsf{T} \mathbf{x} = \hat{\mathbf{V}} \Big( \sum_{k=0}^{K} h_k \hat{\boldsymbol{\Lambda}}^k \Big) \hat{\mathbf{V}}^\mathsf{T} \mathbf{x}$$
Frequency response PCA

#### CoVariance filter

- ightharpoonup Analogy between  $\mathbf{H}(\hat{\mathbf{C}})$  and PCA
  - Using eigendecomposition  $\,\hat{\mathbf{C}} = \hat{\mathbf{V}}\hat{\mathbf{\Lambda}}\hat{\mathbf{V}}^\mathsf{T}\,$  , it follows that

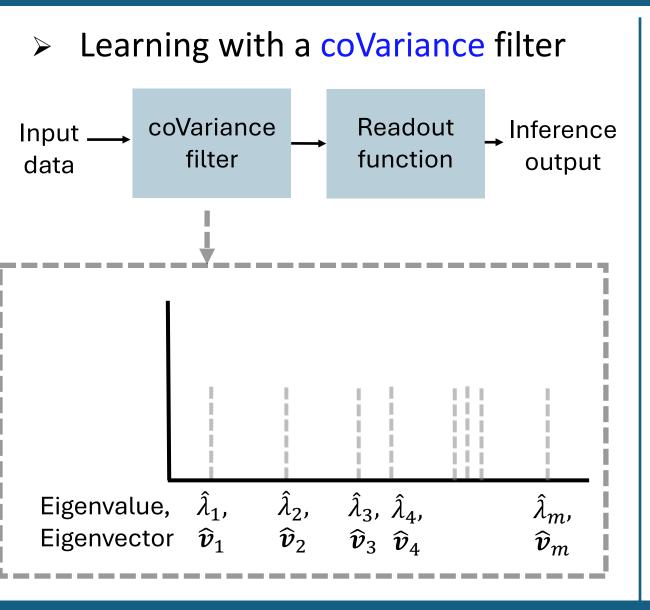
$$\mathbf{z} = \mathbf{H}(\hat{\mathbf{C}})\mathbf{x} = \sum_{k=0}^{K} h_k \hat{\mathbf{C}}^k \mathbf{x} = \sum_{k=0}^{K} h_k \hat{\mathbf{V}} \hat{\boldsymbol{\Lambda}}^k \hat{\mathbf{V}}^\mathsf{T} \mathbf{x} = \hat{\mathbf{V}} \Big( \sum_{k=0}^{K} h_k \hat{\boldsymbol{\Lambda}}^k \Big) \hat{\mathbf{V}}^\mathsf{T} \mathbf{x}$$
Frequency response PCA

GFT of coVariance filter output z and PCA are equivalent

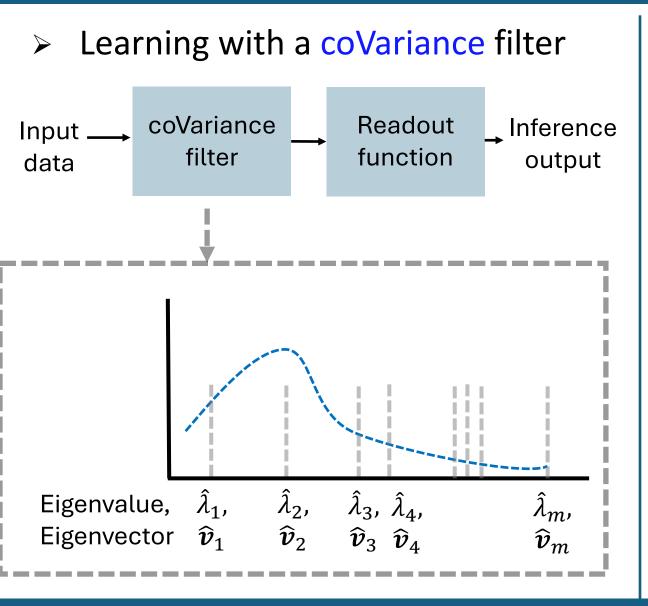
$$\tilde{\mathbf{z}} = \Big(\sum_{k=0}^K h_k \hat{\mathbf{\Lambda}}^k\Big) \hat{\mathbf{V}}^\mathsf{T} \mathbf{x}$$

*i*-th component of  $\tilde{\mathbf{z}}$  is modulated by  $h(\lambda_i) = \sum_{k=0}^K h_k \lambda_i^k$ 

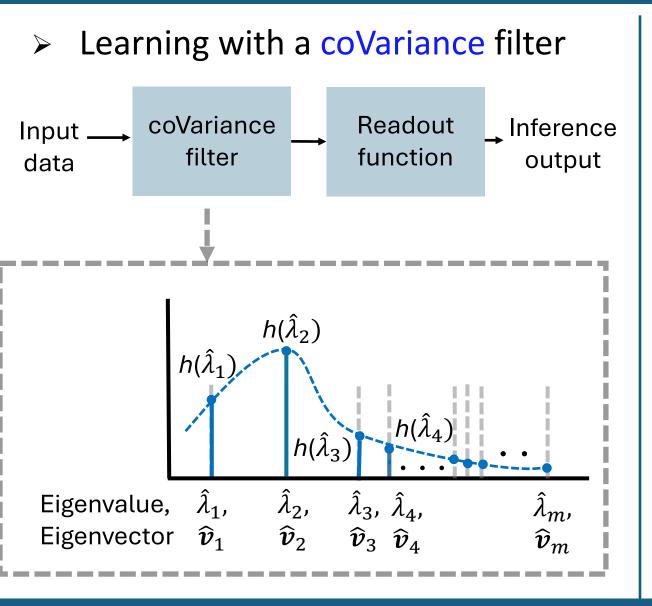
# Learning with coVariance filter versus PCA-based learning



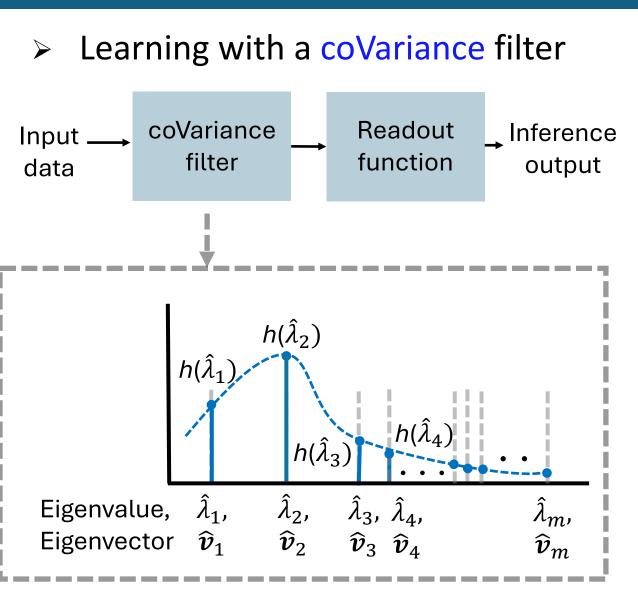
# Learning with coVariance filter versus PCA-based learning

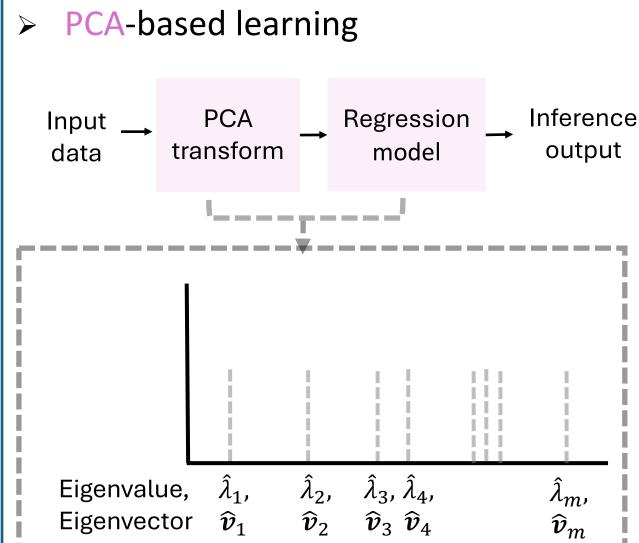


# Learning with coVariance filter versus PCA-based learning

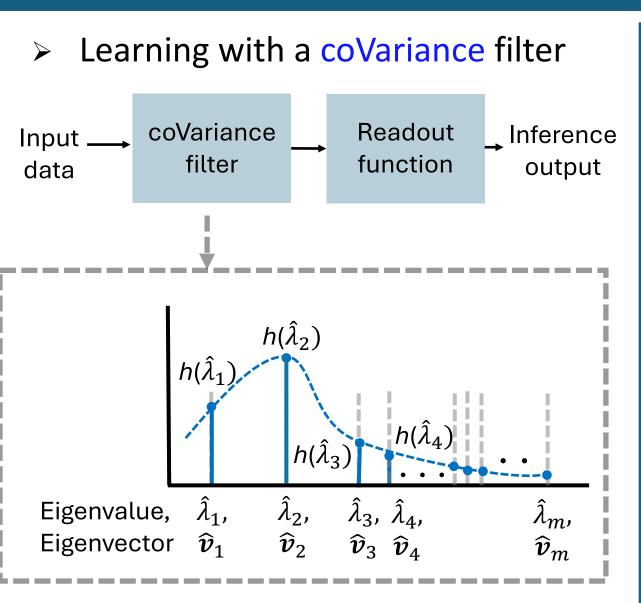


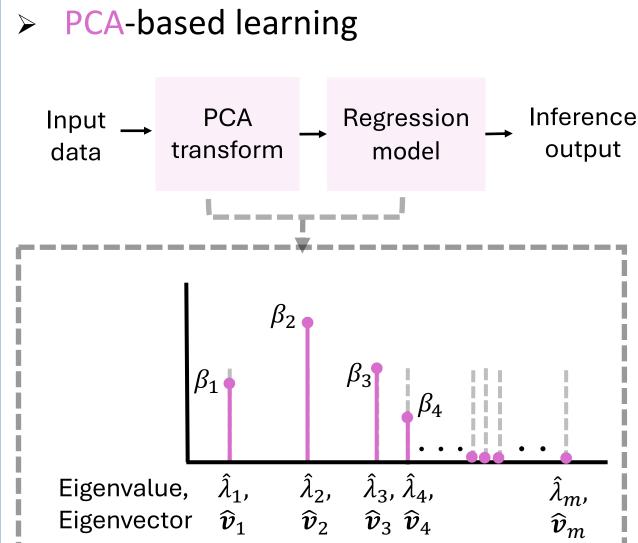
#### Learning with coVariance filter versus PCA-based learning





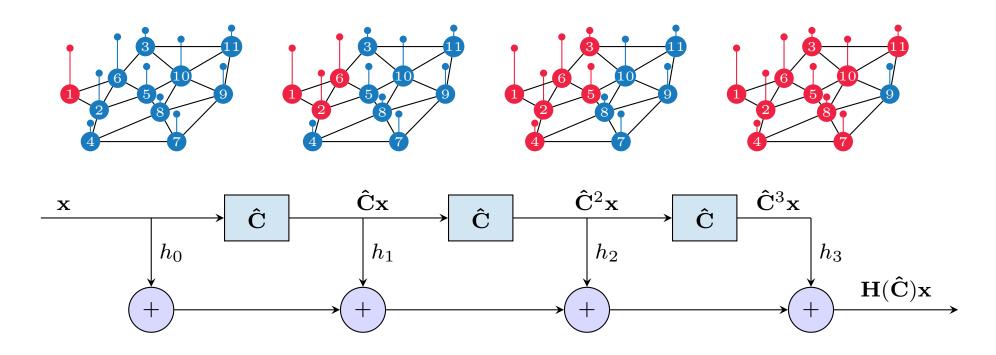
## Learning with coVariance filter versus PCA-based learning





#### coVariance filters as convolutional operators

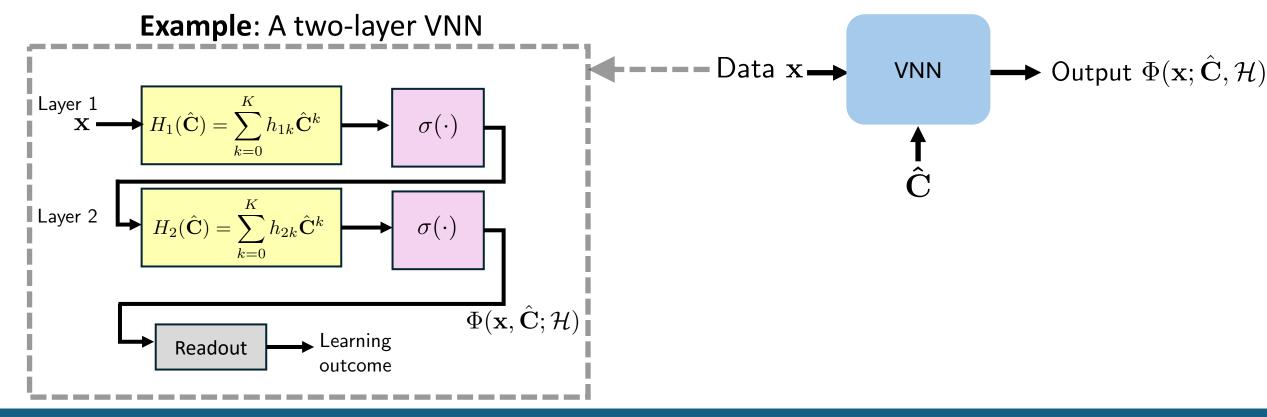
ightharpoonup Operation  $\hat{\mathbf{C}}^k\mathbf{x}$  performs a k-shift of signal  $\mathbf{x}$  over graph defined by  $\hat{\mathbf{C}}$ 



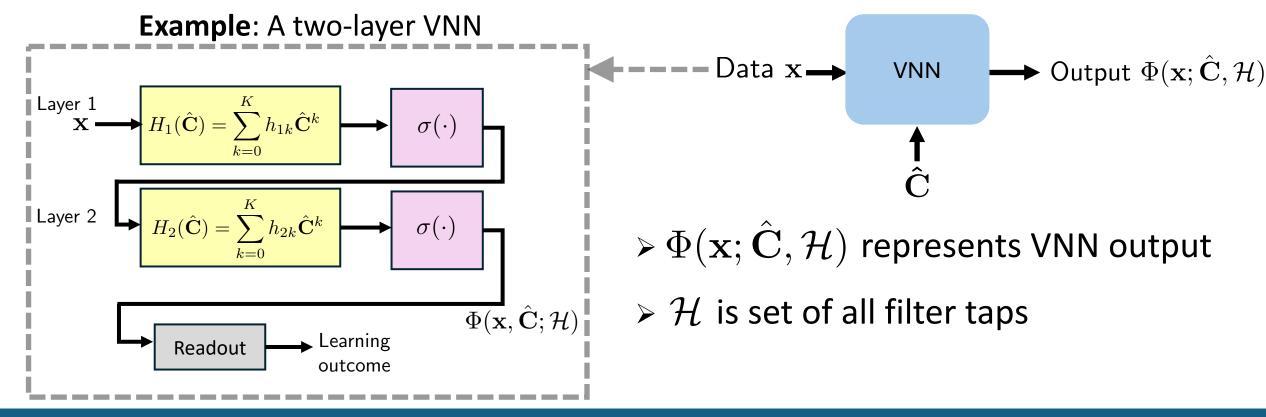
 $\succ$  Parameters  $\{m{h}_{m{k}}\}$  are called **filter taps**, are **scalars** and **learnable** parameters

- > coVariance filters can learn only linear representations
- $\succ$  To accommodate learn **non-linear** representations, concatenate coVariance filter with pointwise non-linearity  $\sigma$  (for e.g., ReLU, sigmoid, etc.)

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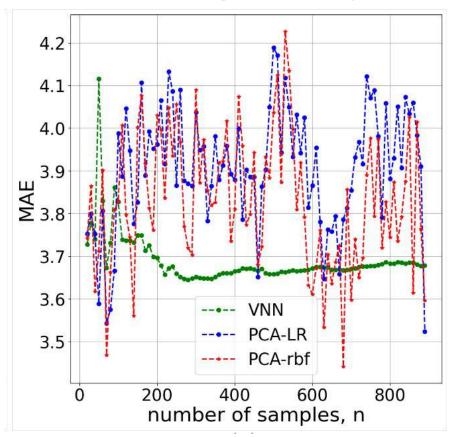
- > coVariance filters can learn only linear representations
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#### VNNs outperform PCA (regression task)

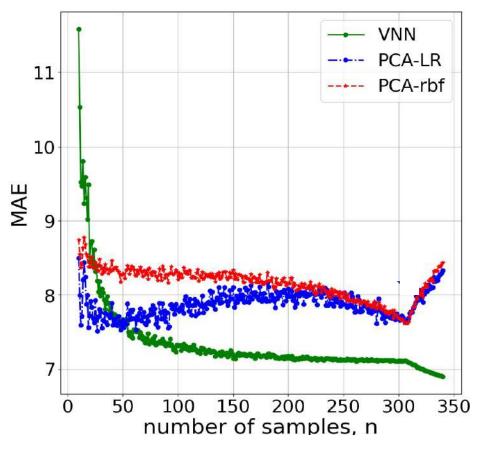
#### Synthetic data

(Friedman regression problem)



#### Neuroimaging data

(age prediction task)



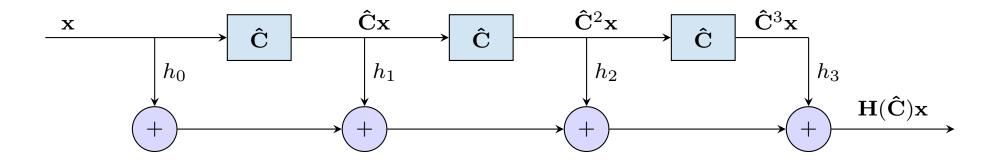
#### Covariance Filters and Neural Networks

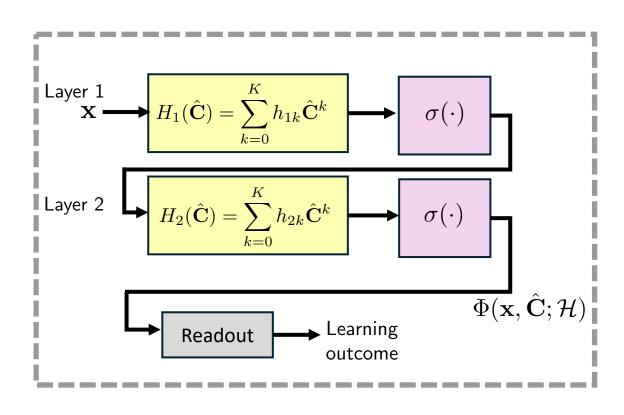
#### Covariance filters

ightharpoonup A covariance filter is a polynomial in the covariance matrix  $\hat{\mathbf{C}}$ 

$$\mathbf{H}(\hat{\mathbf{C}}) = \sum_{k=0}^{K} h_k \hat{\mathbf{C}}^k \mathbf{x}$$

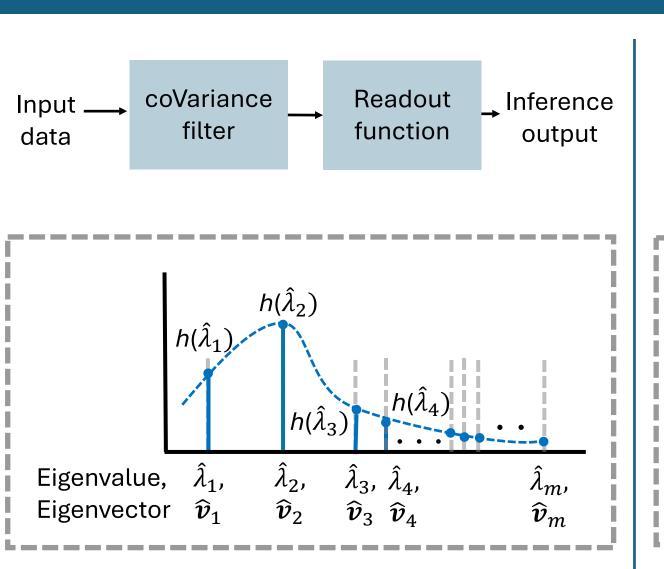
 $\triangleright$  We train the filter coefficients  $h_k$  to accomplish some task

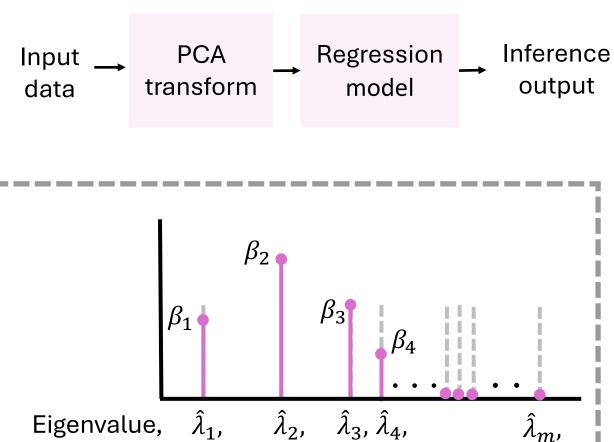




- A VNN is a composition of layers
- > Each of which is a composition of
  - ... a covariance filter
  - ... with a pointwise nonlinearity
- $ightarrow \Phi(\mathbf{x};\hat{\mathbf{C}},\mathcal{H})$  represents VNN output
- $\succ \mathcal{H}$  is the set of trainable filter taps

#### Covariance Filters are Implicitly Equivalent to PCA





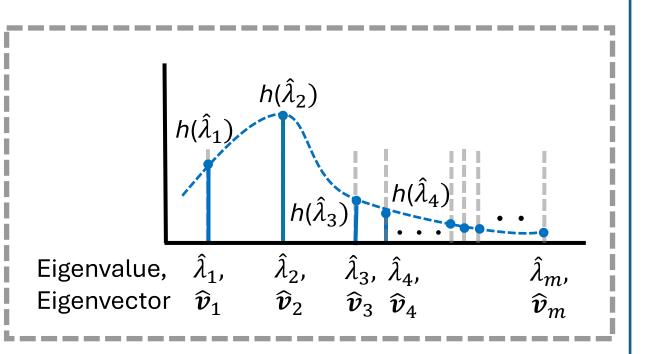
Eigenvector

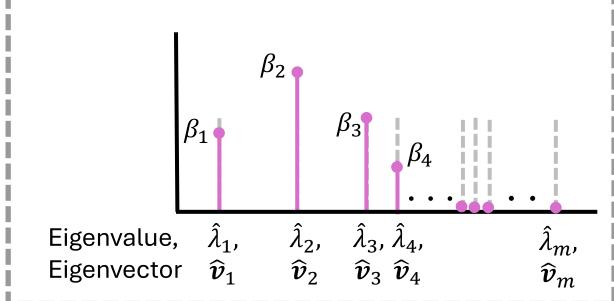
#### Covariance Filters are Implicitly Equivalent to PCA

The difference is that covariance filters (and VNNs) do not require eigenvectors

Stability: Leading to more stable signal processing

**Transferability:** And the possibility of transferring trained filters across scales





#### Stable Inference with VNNs

#### Stability of inference with PCA and VNNs

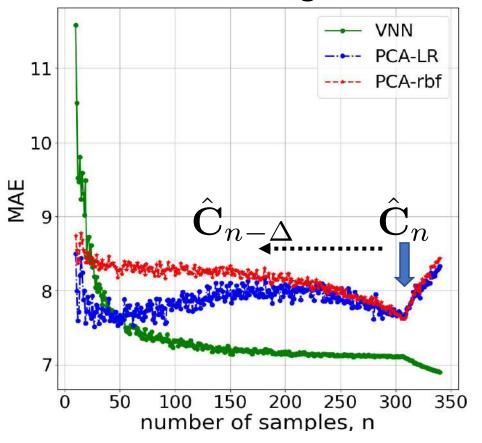
- > PCA-driven inference can be unstable
  - stochastic perturbations due to finite sample effect

VNNs provide stable outcomes

enhanced reproducibility

avoid overfitting

#### Performance on regression task



 $\hat{\mathbf{C}}_n$ : estimated from n samples

# Stochastic perturbations in sample covariance matrix

 $\triangleright$  Recall: Sample covariance matrix  $\hat{\mathbf{C}}$  is estimate of true covariance matrix  $\hat{\mathbf{C}}$ 

$$\hat{\mathbf{C}} = \frac{1}{n-1} \sum_{i=1}^{n} (\mathbf{x}_i - \hat{\boldsymbol{\mu}}) (\mathbf{x}_i - \hat{\boldsymbol{\mu}})^{\mathsf{T}} \qquad \mathbf{C} = \mathbb{E}[(\mathbf{x} - \boldsymbol{\mu}) (\mathbf{x} - \boldsymbol{\mu})^{\mathsf{T}}]$$

 $\Longrightarrow$  eigenvectors/eigenvalues  $\hat{\mathbf{V}}, \hat{\boldsymbol{\Lambda}}$  of  $\hat{\mathbf{C}}$  are estimates of  $\mathbf{V}, \boldsymbol{\Lambda}$  of  $\mathbf{C}$ 

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- ightharpoonup Convergence between  $\hat{\mathbf{V}}, \hat{\mathbf{\Lambda}}$  and  $\mathbf{V}, \mathbf{\Lambda}$  [\*]

$$\|\hat{\mathbf{V}}\mathbf{x} - \mathbf{V}\mathbf{x}\| = \mathcal{O}\left(\frac{1}{n^{1/2}\min_{i \neq j} |\lambda_i - \lambda_j|}\right)$$

## Stochastic perturbations in sample covariance matrix

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- Convergence between  $\hat{\mathbf{V}}$ ,  $\hat{\mathbf{\Lambda}}$  and  $\mathbf{V}$ ,  $\mathbf{\Lambda}$  [\*]

$$\|\hat{\mathbf{V}}\mathbf{x} - \mathbf{V}\mathbf{x}\| = \mathcal{O}\left(\frac{1}{n^{1/2}\min_{i \neq j} |\lambda_i - \lambda_j|}\right)$$



Unstable PCA transform when eigenvalues of covariance are close

[\*] Loukas, Andreas, 2017

How to gauge stability?

$$\mathbf{x} \longrightarrow \mathbf{H}(\hat{\mathbf{C}}) \longrightarrow \mathbf{z} = \mathbf{H}(\hat{\mathbf{C}})\mathbf{x} \qquad \hat{\mathbf{C}} = \frac{1}{n-1} \sum_{i=1}^{n} (\mathbf{x}_i - \hat{\boldsymbol{\mu}})(\mathbf{x}_i - \hat{\boldsymbol{\mu}})^{\mathsf{T}}$$

 $\implies$  Output **z** must be robust to number of samples n used to estimate  $\hat{\mathbf{C}}$ 

How to gauge stability?

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- $\implies$  Output **z** must be robust to number of samples n used to estimate  $\hat{\mathbf{C}}$
- Compare filter outputs for sample and true covariance matrix

$$\mathbf{x} \longrightarrow \mathbf{H}(\hat{\mathbf{C}})$$
  $\longrightarrow \mathbf{z} = \mathbf{H}(\hat{\mathbf{C}})\mathbf{x}$   $\mathbf{x} \longrightarrow \mathbf{H}(\mathbf{C})$   $\longrightarrow \mathbf{z} = \mathbf{H}(\mathbf{C})\mathbf{x}$ 

 $\implies$  metric of interest:  $\|\mathbf{H}(\hat{\mathbf{C}}) - \mathbf{H}(\mathbf{C})\|$ 

$$\mathbf{x} \longrightarrow \mathbf{H}(\hat{\mathbf{C}}) \longrightarrow \mathbf{z} = \mathbf{H}(\hat{\mathbf{C}})\mathbf{x} \qquad \mathbf{x} \longrightarrow \mathbf{H}(\mathbf{C}) \longrightarrow \mathbf{z} = \mathbf{H}(\mathbf{C})\mathbf{x}$$

Stability result [Sihag et al., 2022]

$$\left\| \mathbf{H}(\hat{\mathbf{C}}) - \mathbf{H}(\mathbf{C}) \right\| = \mathcal{O}\left(\frac{1}{n^{1/2 - \varepsilon}}\right)$$

coVariance filter output is asymptotically consistent

$$\mathbf{z} \longrightarrow \mathbf{H}(\hat{\mathbf{C}}) \longrightarrow \mathbf{z} = \mathbf{H}(\hat{\mathbf{C}})\mathbf{x} \qquad \mathbf{x} \longrightarrow \mathbf{H}(\mathbf{C}) \longrightarrow \mathbf{z} = \mathbf{H}(\mathbf{C})\mathbf{x}$$

Stability result [Sihag et al., 2022]

$$\left\| \mathbf{H}(\hat{\mathbf{C}}) - \mathbf{H}(\mathbf{C}) \right\| = \mathcal{O}\left(\frac{1}{n^{1/2 - \varepsilon}}\right)$$

Assumption.

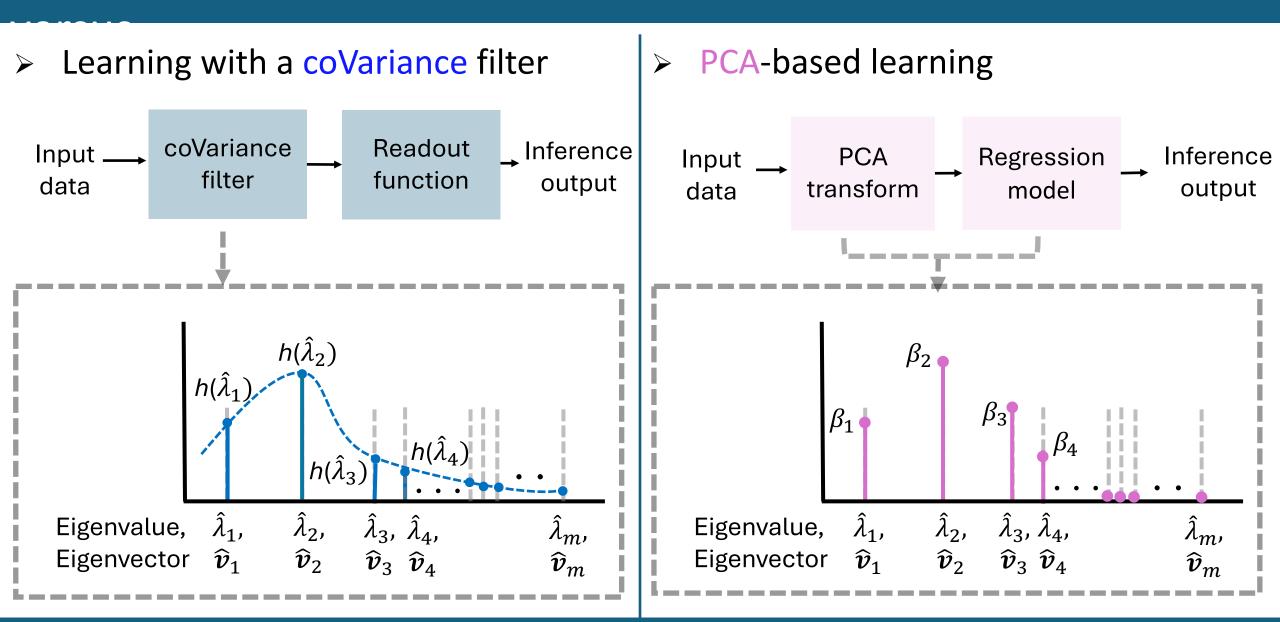
Frequency response of filter  $\mathbf{H}(\mathbf{C})$  satisfies

$$|h(\lambda_i) - h(\lambda_j)| \le Q \frac{|\lambda_i - \lambda_j|}{k_i}$$

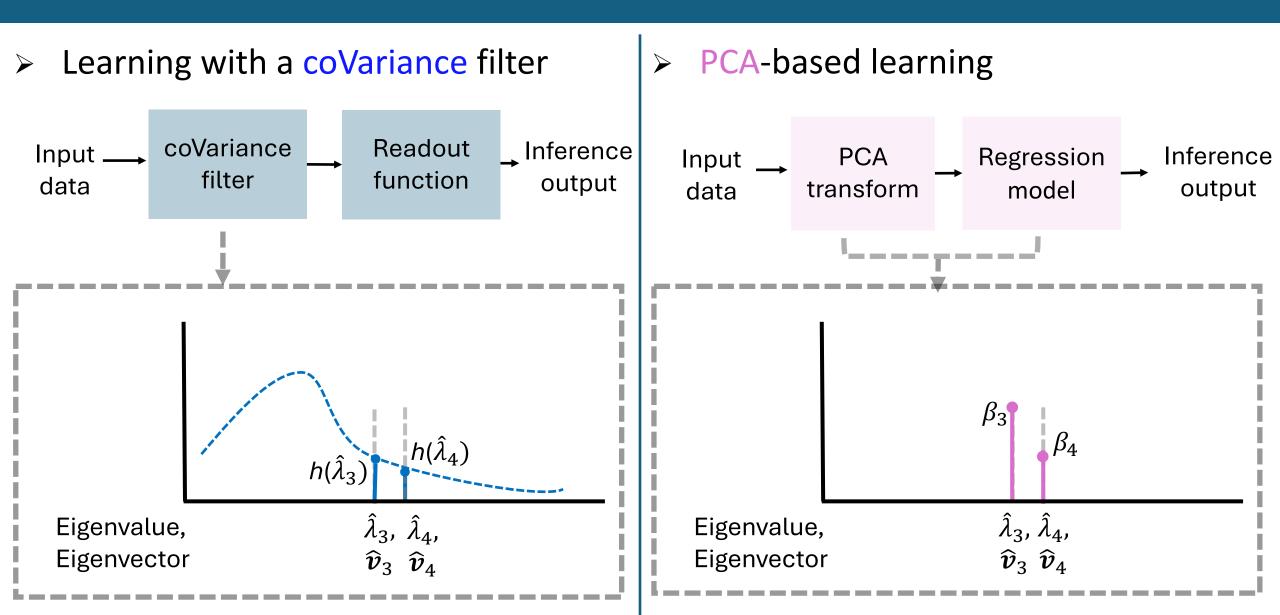
coVariance filter output is asymptotically consistent

coVariance filter sacrifices discriminability between close eigenvalues for stability

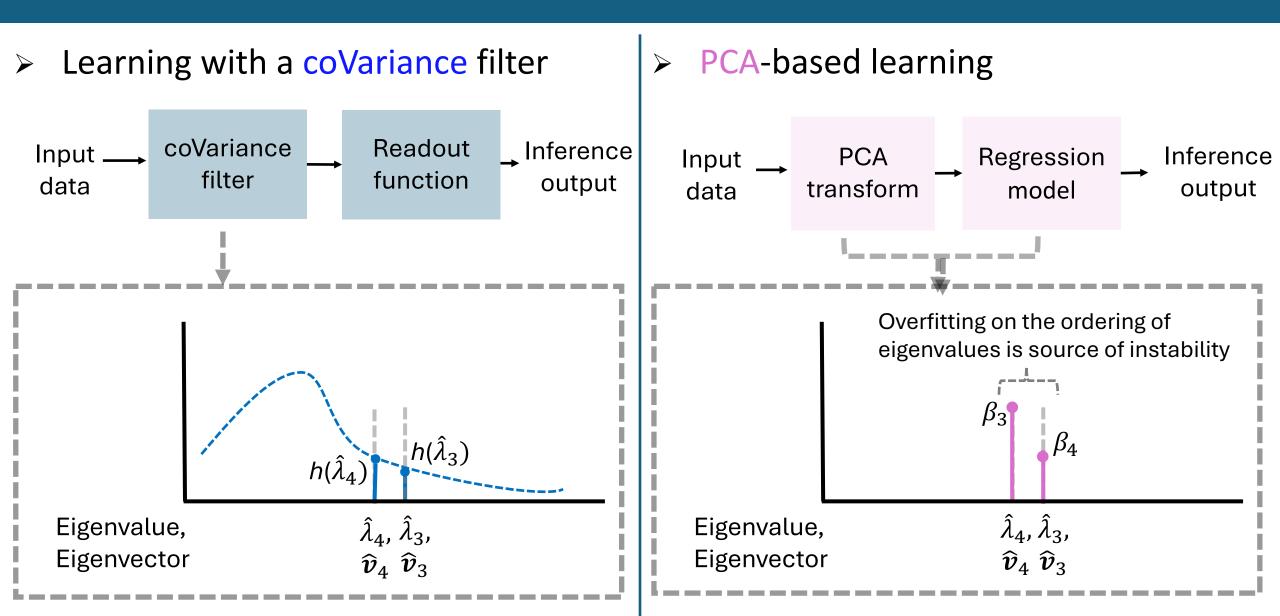
#### Recall: Learning with coVariance filter versus PCA-based learning



# Why is coVariance filter more stable than PCA?



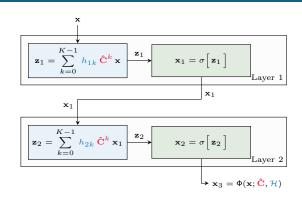
# Why is coVariance filter more stable than PCA?



# Stability of VNNs

- VNNs inherit the stability from coVariance filters
  - Stability bound depends on the bound for filters

$$\|\mathbf{H}(\hat{\mathbf{C}}) - \mathbf{H}(\mathbf{C})\| = \mathcal{O}\left(\frac{1}{n^{\frac{1}{2}} - \varepsilon}\right) = \alpha_n$$



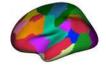
For a VNN with L layers and F filters in parallel,

$$\left\| \Phi(\mathbf{x}, \hat{\mathbf{C}}; \mathcal{H}) - \Phi(\mathbf{x}, \mathbf{C}; \mathcal{H}) \right\| \leq LF^{L-1} \alpha_n$$

Stability bound increases with number of layers and size of filter banks

#### Stability of VNNs: Experiments

Regression task

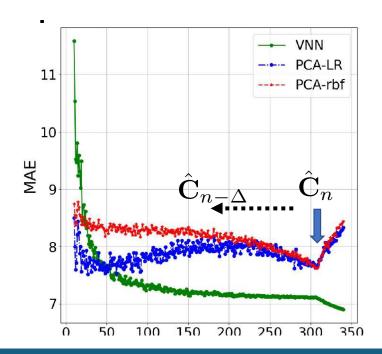


Cortical thickness — VNN — Estimate of age

Comparison against PCA-regression

**Data**: cortical thickness dataset (m = 104) from (n = 341) human subjects

Metric: MAE (mean absolute error)



VNN: coVariance Neural Network

PCA-LR: PCA-regression with linear kernel

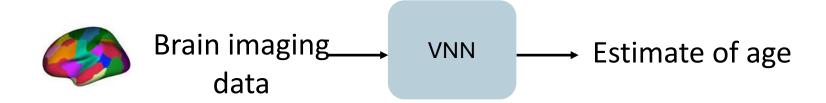
PCA-rbf: PCA regression with rbf kernel

VNN outperforms PCA and is more stable

#### Transferability of VNNs

## Empirical evidence of transferability across multiscale data

- > Transferability across multiscale datasets
  - Multiscale datasets capture same phenomenon at different scales



Transferability across datasets with different number of features

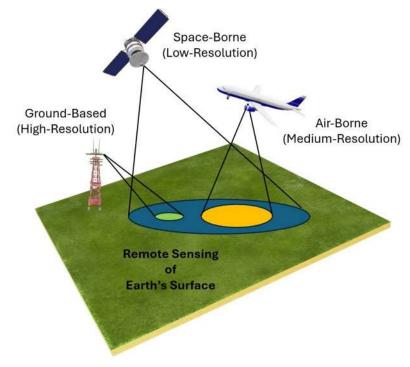
Testing		
Training	100-feature dataset	300-feature dataset
100-feature dataset	$5.39 \pm 0.084$	$5.5 \pm 0.101$

## Transferability

- > Learning models could generalize to compatible datasets
- > compatible: different dimensionalities and describing the same domain

Brain imaging data

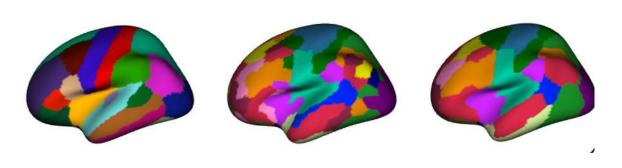
#### Remote sensing



Credit: Mustafa Aksoy, UAlbany

#### Transferability

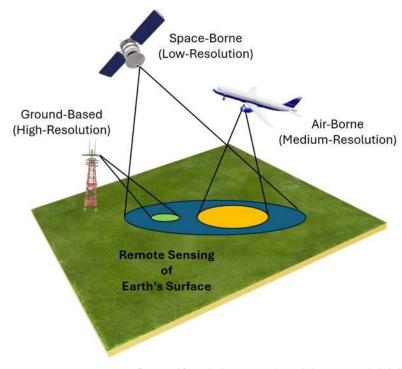
- > Learning models could generalize to compatible datasets
- compatible: different dimensionalities and describing the same domain



Brain imaging data

Motivation: novel metric for generalizability, managing high dimensional data...

#### Remote sensing

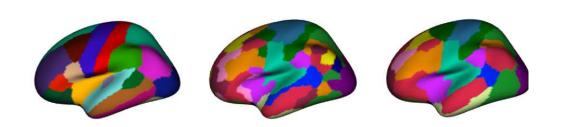


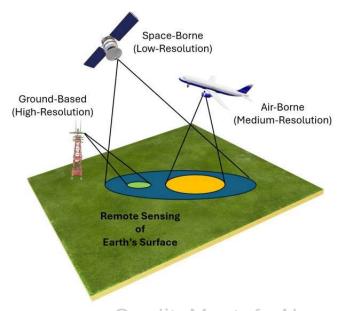
Credit: Mustafa Aksoy, UAlbany

## Transferability

- > Most statistical approaches, including PCA, operate within the dimensionality
  - ightharpoonup > seamless transference not possible across different dimensionalities
- This section: How do VNNs transfer?

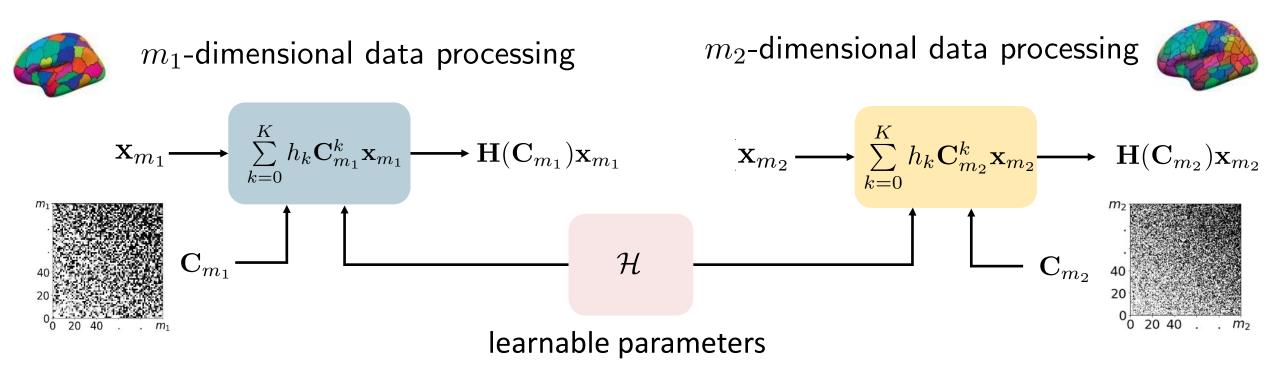
When is transference successful?





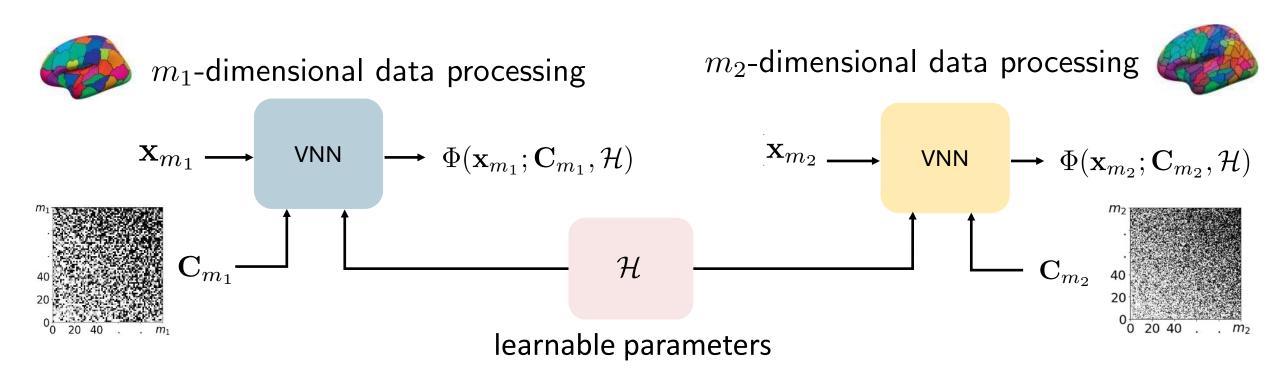
Credit: Mustafa Aksoy, UAlbany

#### coVariance filters are scale-free models



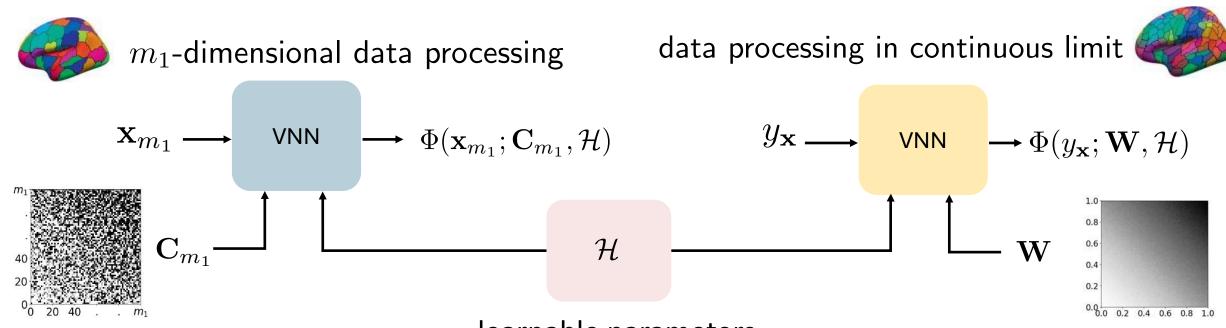
 $\blacktriangleright$  A coVariance filter  $\mathbf{H}(\cdot)$  with scalar filter taps  $\{h_k\}$  can process dataset (covariance matrix) of any arbitrary dimensionality: **scale-free model** 

#### VNNs as scale-free models



How to compare  $\Phi(\mathbf{x}_{m_1}; \mathbf{C}_{m_1}, \mathcal{H})$  and  $\Phi(\mathbf{x}_{m_2}; \mathbf{C}_{m_2}, \mathcal{H})$ ?

#### VNNs as scale-free models



learnable parameters

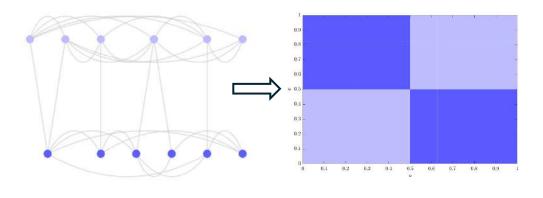
Continuous limit of covariance matrices as  $m \to \infty$ 

How to compare  $\Phi(\mathbf{x}_{m_1}; \mathbf{C}_{m_1}, \mathcal{H})$  and  $\Phi(y_{\mathbf{x}}; \mathbf{W}, \mathcal{H})$ ?

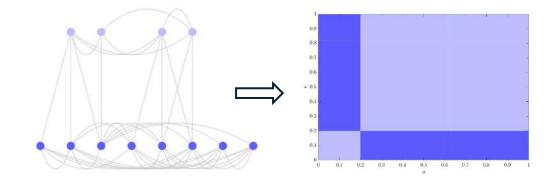
# Graphons as continuous limits

> Graphs can have **limit objects** with uncountable number of nodes

> Example: Stochastic block models [Ruiz et al., 2021]



**Balanced SBM** 



**Unbalanced SBM** 

# Graphons as continuous limits

- > Graphon: A graphon is a symmetric, bounded measurable function
  - Node labels are graphon arguments  $u \in [0,1]$
  - edge weights are graphon values  $\mathbf{W}(u, v) = \mathbf{W}(v, u)$

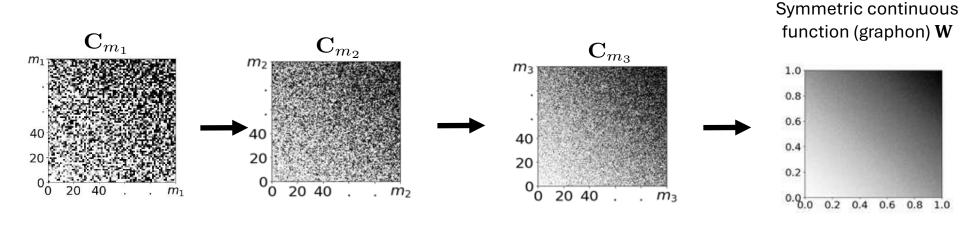
$$\mathbf{W}:[0,1]^2\mapsto \mathbb{R}$$

# Graphons as continuous limits

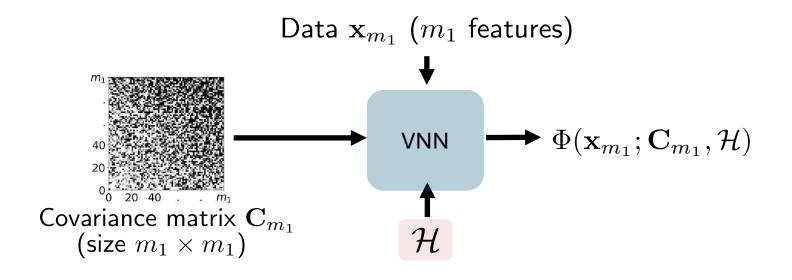
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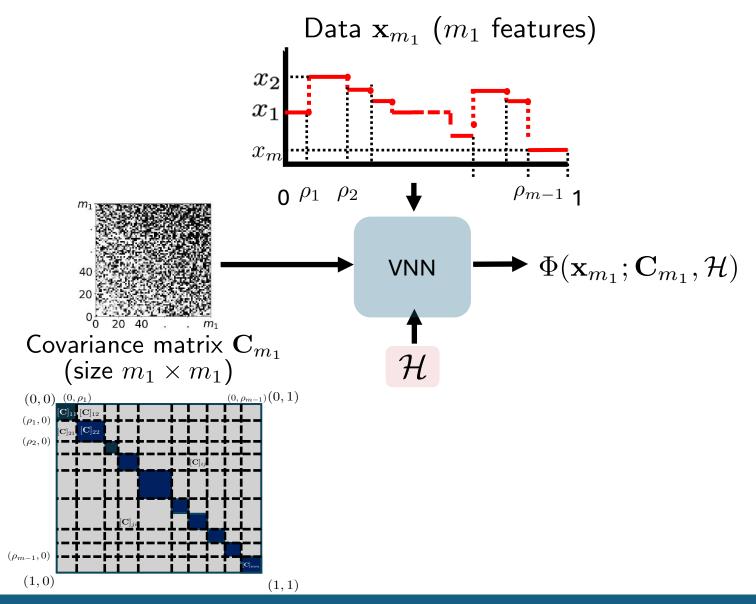
Transferability when covariance matrix is part of some converging sequence



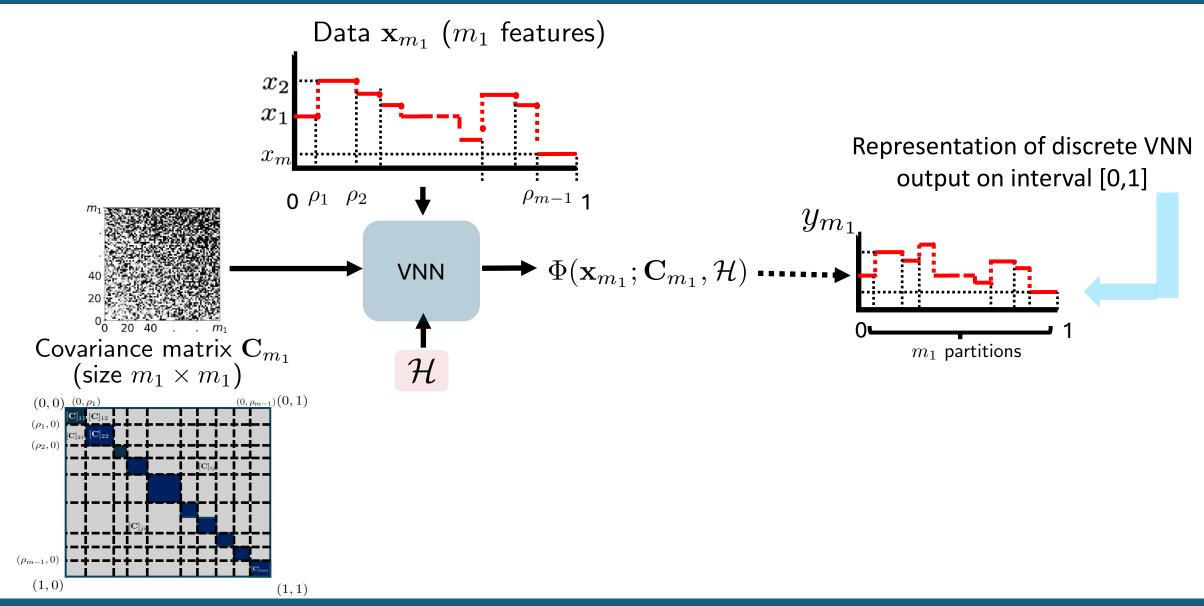
# Redefining VNNs in continuous domain



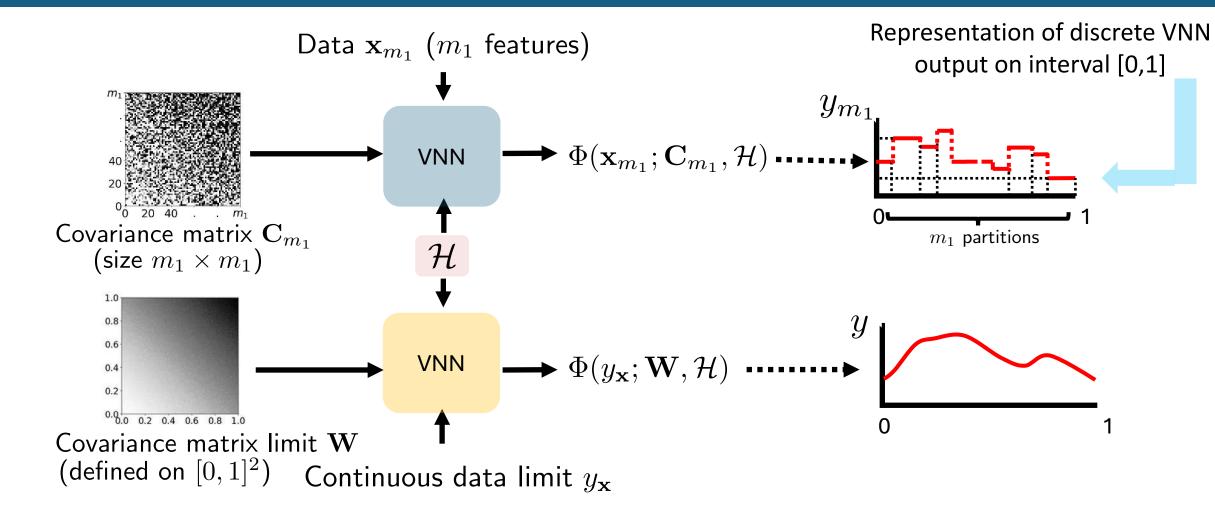
## Redefining VNNs in continuous domain



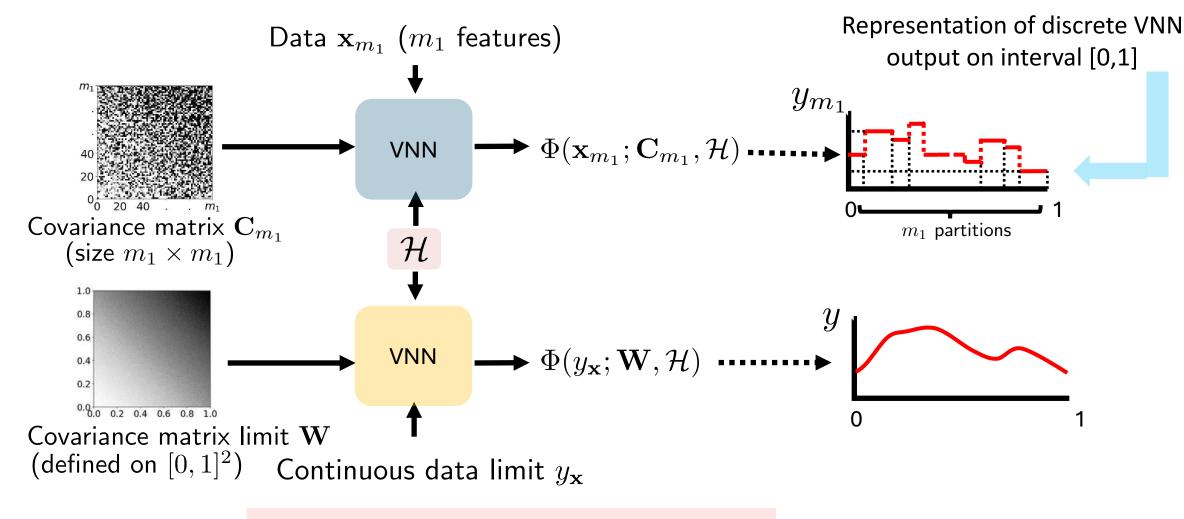
# Redefining VNNs in continuous domain



## Problem formulation for transferability

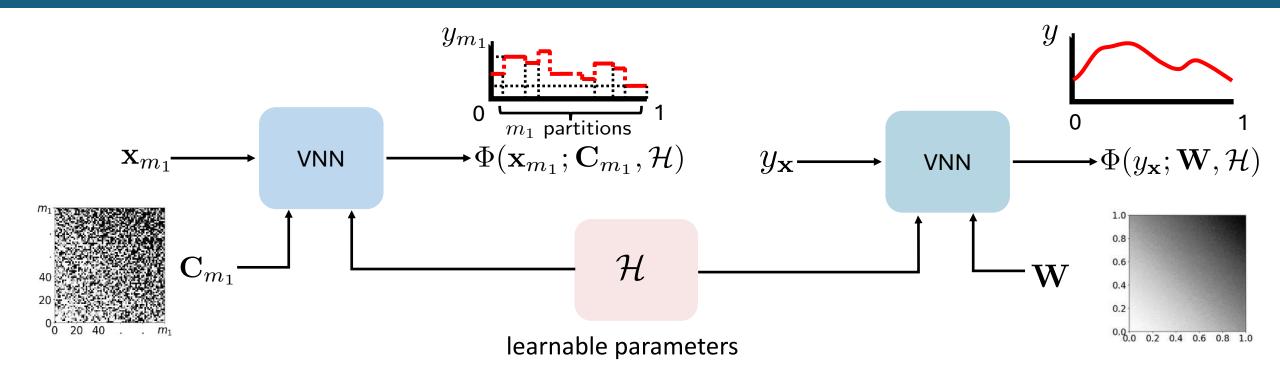


# Problem formulation for transferability



Find  $\vartheta$ , such that,  $||y_{m_1} - y||_2 \leq \vartheta$ 

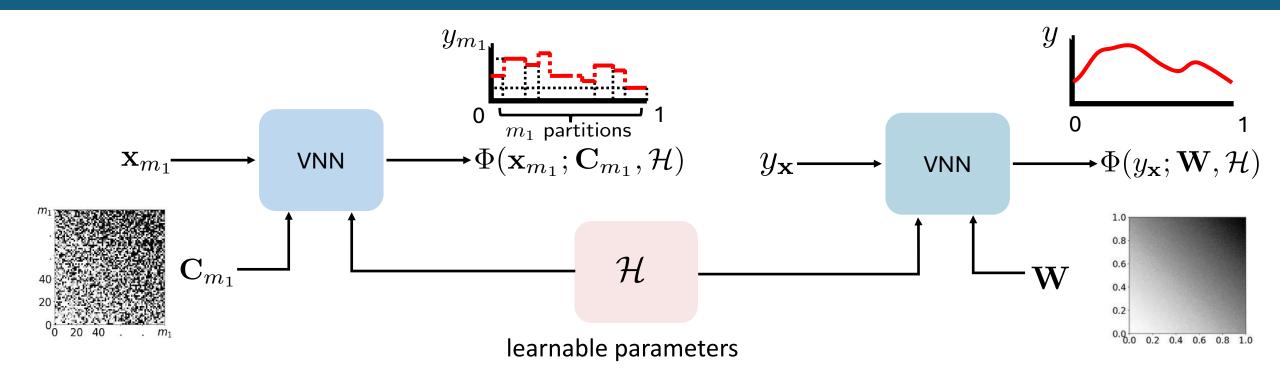
## VNNs are provably transferable

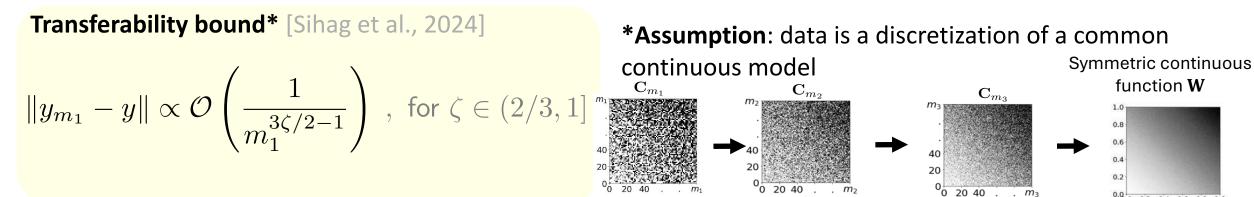


Transferability bound\* [Sihag et al., 2024]

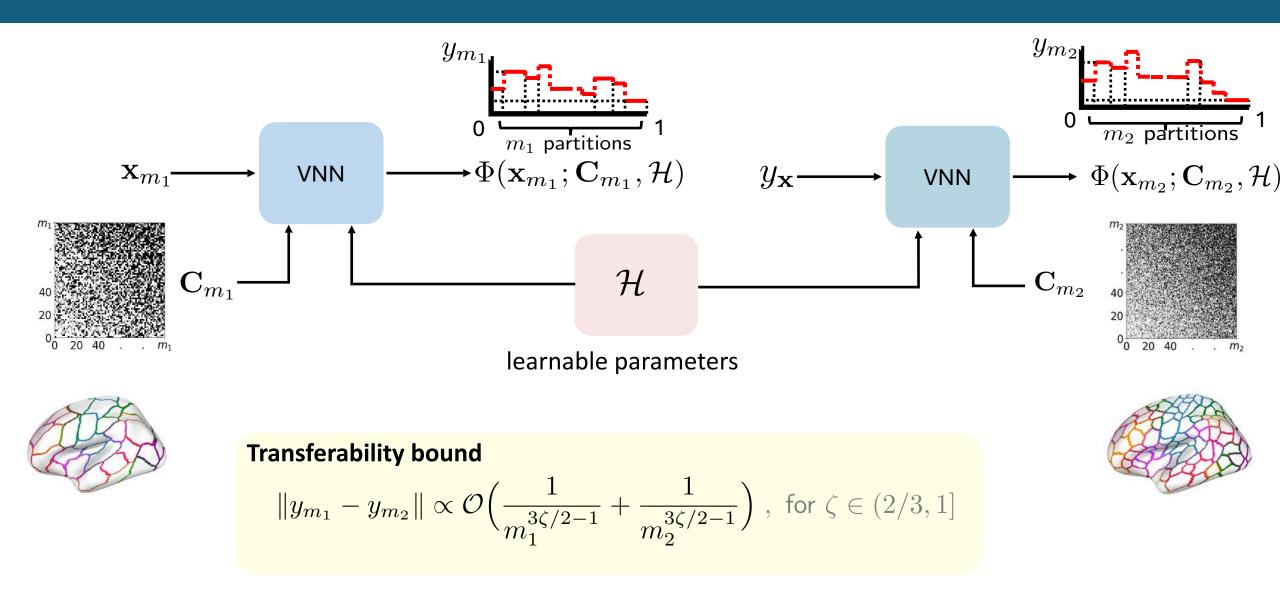
$$||y_{m_1} - y|| \propto \mathcal{O}\left(\frac{1}{m_1^{3\zeta/2-1}}\right), \text{ for } \zeta \in (2/3, 1]$$

## VNNs are provably transferable





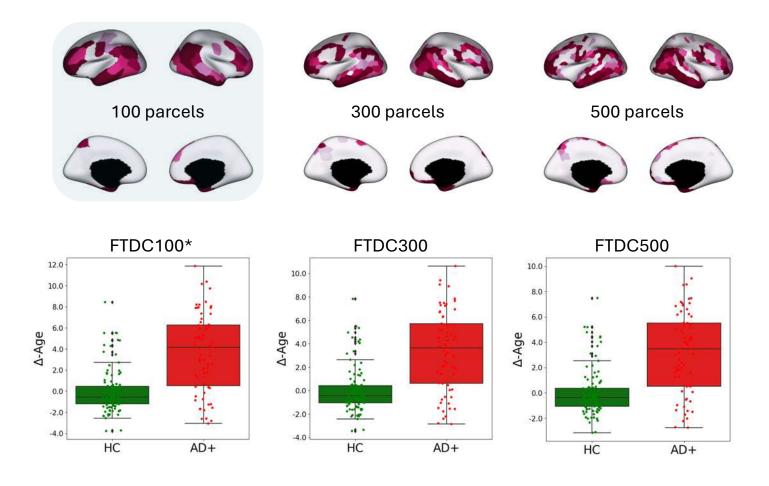
## VNNs are provably transferable



## Experiments

Objective: Brain age gap prediction in HC (healthy) and AD+ (Alzheimer's) cohorts from

VNNs trained on 100-feature dataset [Sihag et al., NeurIPS, 2024, JSTSP 2024, SPM 2025]



- ROIs contributing to elevated brain age gap in AD+ across different resolutions
- Brain age gap is elevated in AD+ w.r.t HC cohort in 100feature dataset
- Results on brain age gap retained after transferring VNN to 300 and 500-feature datasets

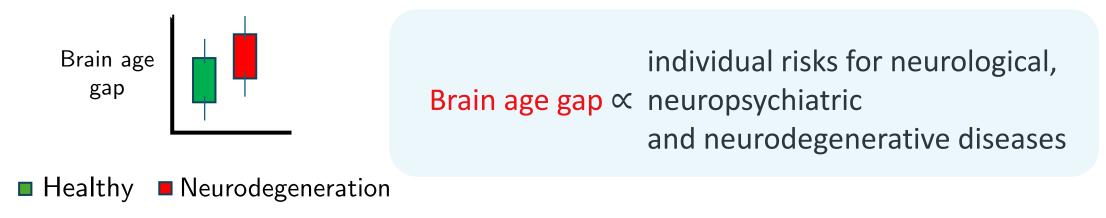
Principled brain age gap prediction with VNNs

# Brain age gap

- > Individual rate of "aging" is different from chronological rate of aging
  - Driven by environment, genetics, neurodegeneration
- > Brain age provides a biological estimate brain age, derived from neuroimaging

## Brain age gap

- > Individual rate of "aging" is different from chronological rate of aging
  - Driven by environment, genetics, neurodegeneration
- > Brain age provides a biological estimate brain age, derived from neuroimaging
- > The brain age gap is the deviation between brain age and chronological age

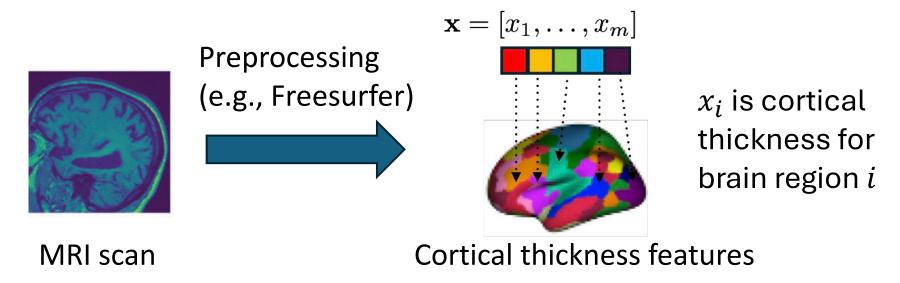


# Neurodegeneration (in terms of cortical atrophy)

- > Neurodegeneration is accelerated decline of structure or function of the brain
- > Cortical atrophy: reduction in cortical thickness/volume/area

(characteristic of healthy aging and disorders like Alzheimer's disease (AD),

frontotemporal dementia (FTD), etc.)

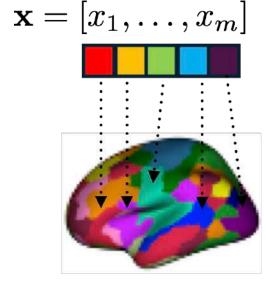


# Neuroimaging Data: Basics

Data sample corresponds to measurement associated with brain (cortical) surface



datasets may have distinct dimensionalities

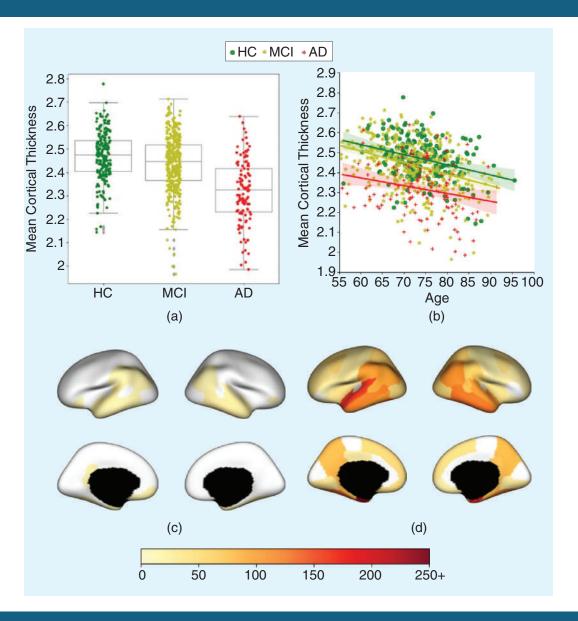


Anatomic features

 Multi-resolution brain atlas discretizes brain surface at multiple resolutions (for e.g., Schaefer's atlas has resolutions 100-1000)

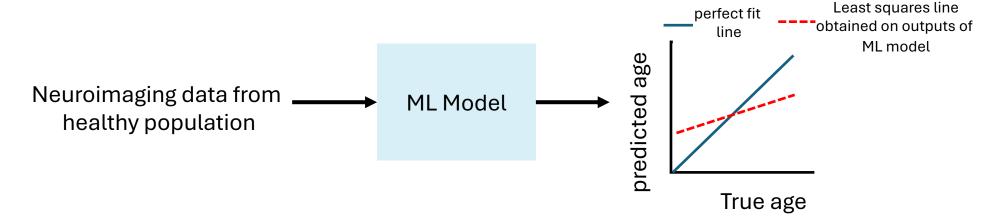
# Case study (Neurodegeneration)

- > **Data:** cortical thickness from 3 cohorts
  - HC (healthy)
  - MCI (Mild cognitive impairment )
  - AD (Alzheimer's disease)
- Larger cortical atrophy is feature of AD
- MCI is precursor to AD
   shows intermediate cortical
   atrophy between HC and AD
- Aging also leads to cortical atrophy



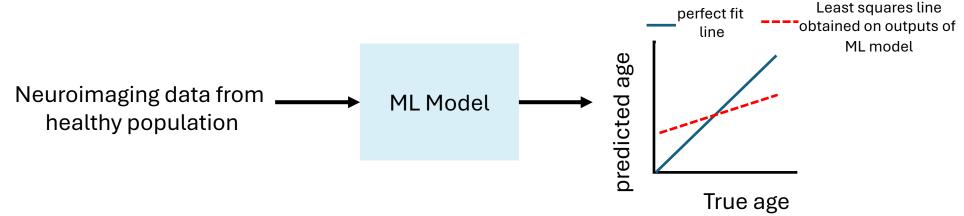
# Brain age gap evaluation using ML

**Step 1.** Train ML model to predict chronological age for healthy controls from cortical thickness features



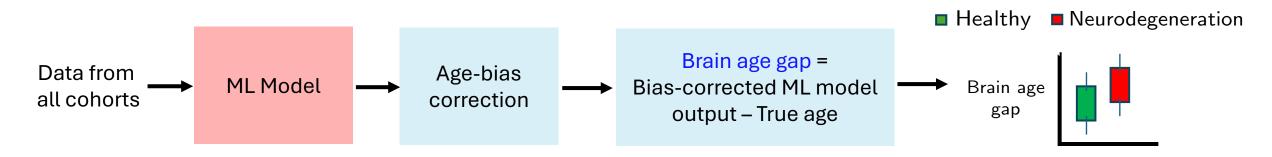
# Brain age gap evaluation using ML

**Step 1.** Train ML model to predict chronological age for healthy controls from cortical thickness features



Step 2. Linear regression-based age-bias correct for outputs of ML model

Step 3. Obtain brain age gap for healthy controls and individuals with neurodegenerative condition.



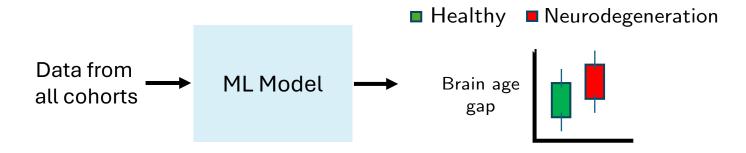
# Brain age gap prediction is a transfer learning problem

Train ML model to predict age on a large dataset (healthy population)

#### **Pre-training**



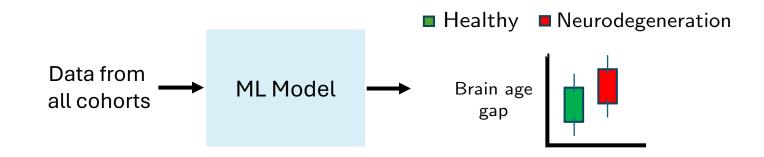
Apply the pre-trained ML model on a target dataset (neurodegeneration)



Brain age gap is the residual of the model

# Brain age gap prediction is a transfer learning problem

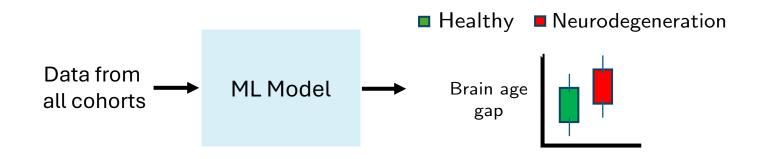
- > Some observations about a meaningful brain age gap
  - We expect model performance to degrade in target population
    - ✓ Degradation in performance (residuals) must be in a specific direction



# Choice of learning parametrization

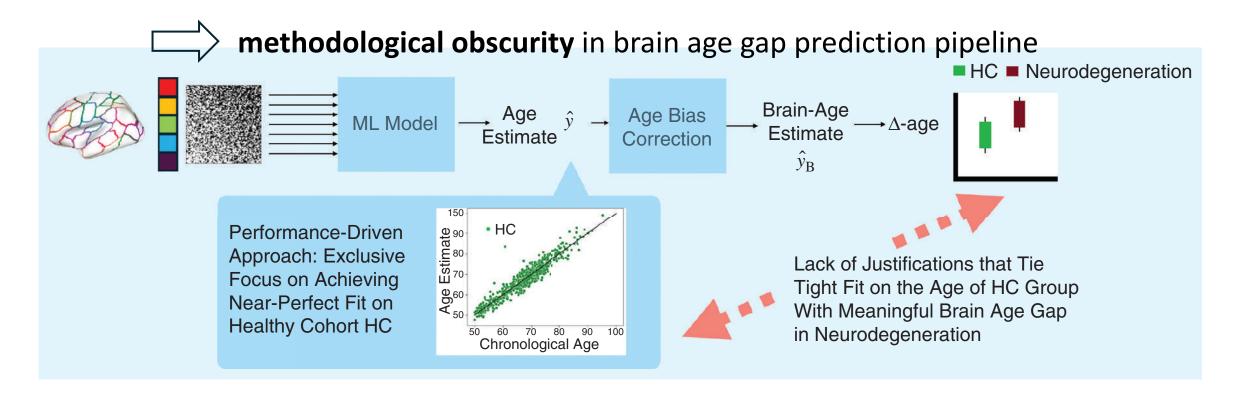
> Choice of ML model dictates how data is leveraged to gauge brain age gap

- > Prevalent approaches focus on achieving perfect pre-training performance
  - Performance-driven approaches
- > Performance-driven approaches do not guarantee `meaningful' brain age gap



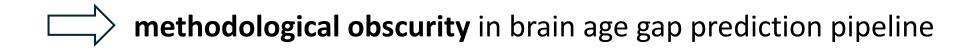
# Choice of learning parametrization

- > Neural networks are prevalent in performance-driven approaches
- A Neural Network may not be interpretable and prone to overfitting



# Choice of learning parametrization

- > Neural networks are prevalent in performance-driven approaches
- A Neural Network may not be interpretable and prone to overfitting



Performance in pre-training does not dictate **meaningful residuals** in target population

# A principled approach to brain age gap prediction

- > Focus on residuals of the ML model, not prediction performance
- > Qualitative evaluation during pre-training

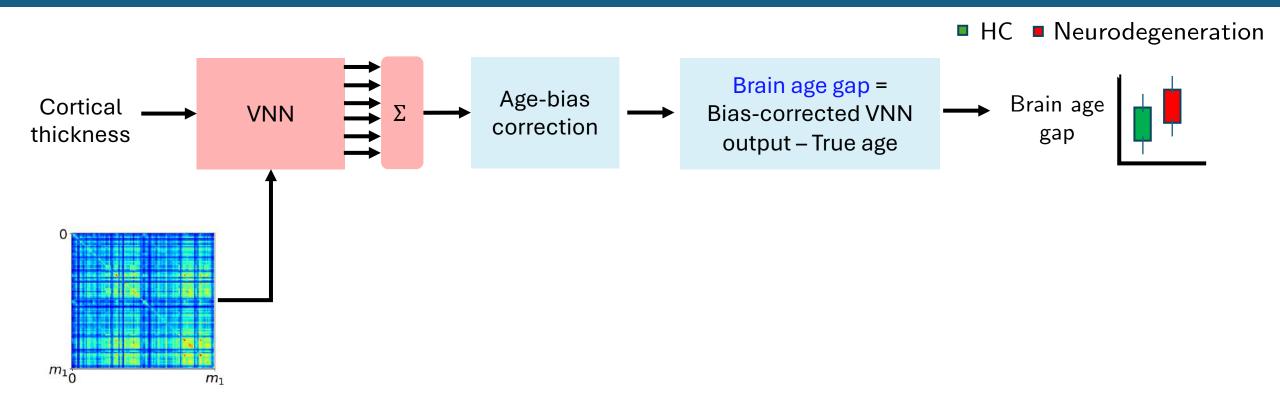
what does the model learn during pre-training on healthy population?

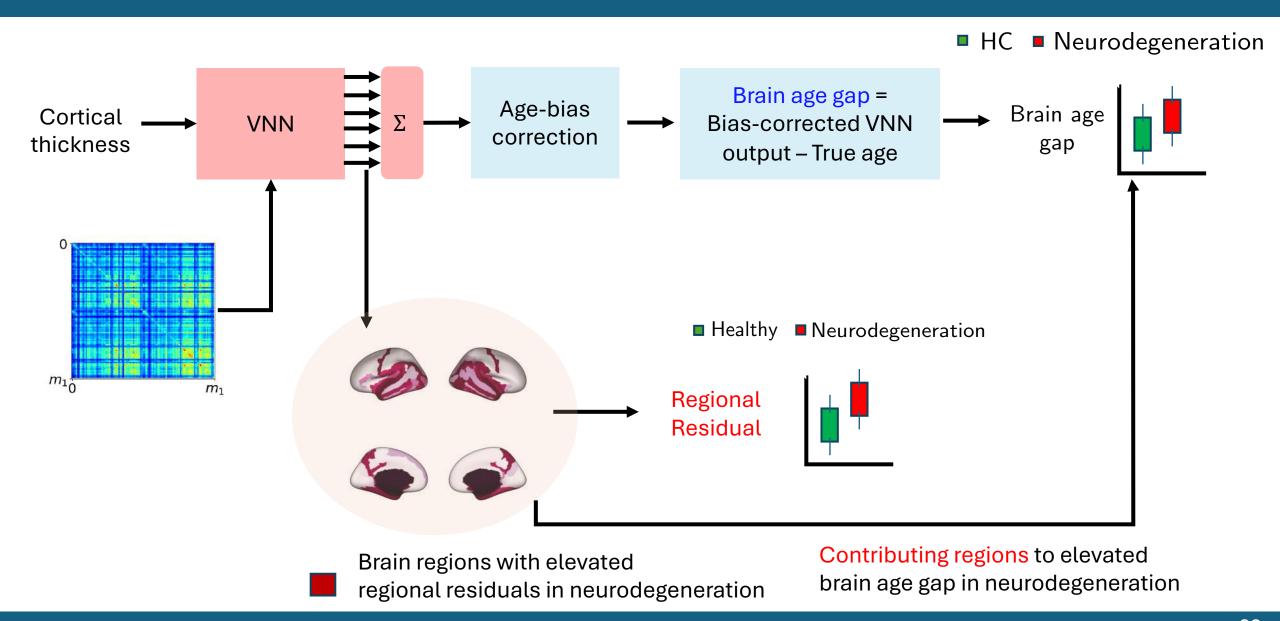
> Interpretability/explainability:

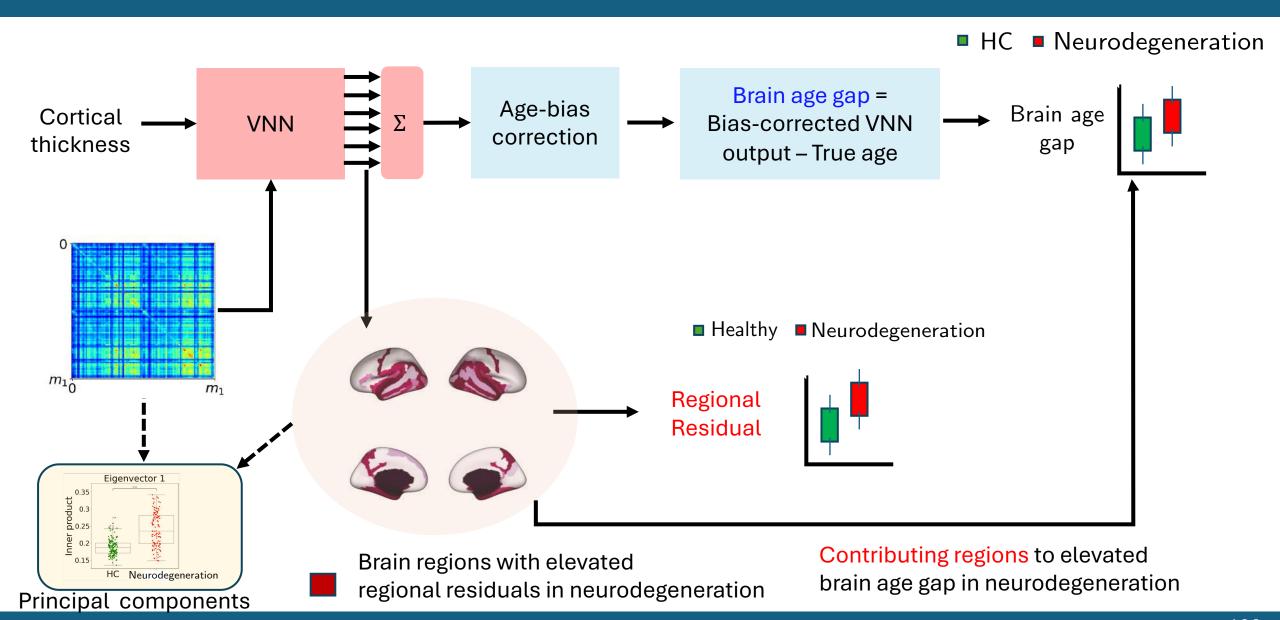
what's driving elevated brain age gap (residuals) in neurodegeneration?

> Generalizability to diverse target populations

Sihag et al., 2025 (SPM, to appear)

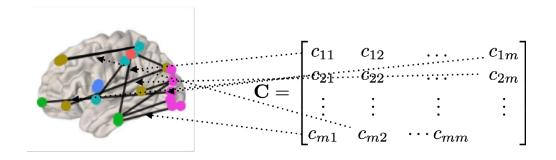




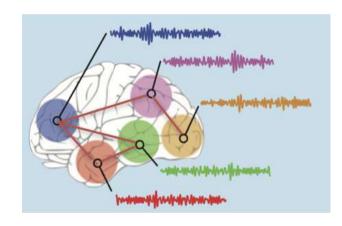


#### Network neuroscience

Modeling brain as a network (connectomes)



Anatomical covariance matrix (structural connectome)



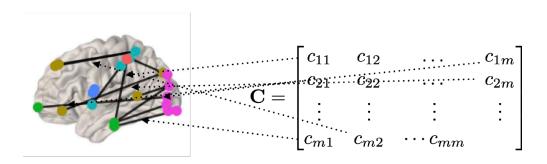
Functional connectome

#### > Motivation

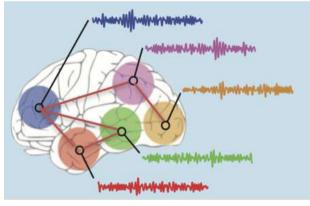
- Significant redundancies in brain structural/functional features
- Brain structure/function is compromised in neurodegeneration

#### Covariance matrices in network neuroscience

> Covariance matrices appear commonly in network neuroscience



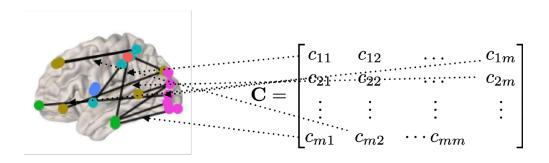
Anatomical covariance matrix



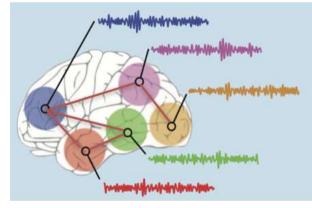
Functional connectome

#### Covariance matrices in network neuroscience

> Covariance matrices appear commonly in network neuroscience

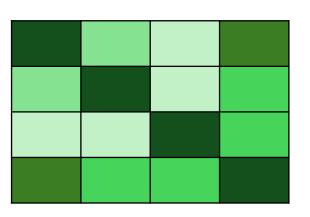


Anatomical covariance matrix



Functional connectome

- > Inference over covariance matrices in network neuroscience
  - Traditional statistical approaches (for e.g., PCA)
    - o Interpretable, suitable for low data regimes
  - Deep learning approaches (for e.g., GNNs)
    - Enhanced expressivity, improved performance



# VNNs are well suited for neuroimaging data analysis

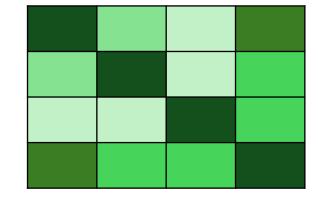
> Properties of VNNs make them appealing for neuroimaging data analysis

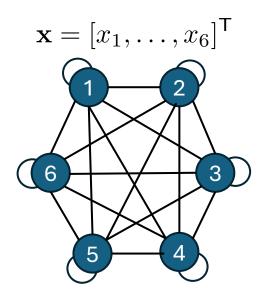
- Connections with PCA transparent outcomes by leveraging spectrum of covariance matrix
- Stability 
   reproducible outcomes in limited data settings

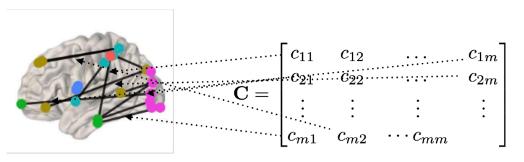
Transferability 
 — enhanced generalizability and robustness to choice of brain atlases

# Anatomical covariance matrix as a graph

Covariance matrix is a data-driven graph







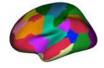
Covariance matrix as a fully-connected graph

$$\hat{\mathbf{C}} = \frac{1}{n-1} \sum_{i=1}^{n} (\mathbf{x}_i - \hat{\boldsymbol{\mu}}) (\mathbf{x}_i - \hat{\boldsymbol{\mu}})^{\mathsf{T}}, \text{ where } \hat{\boldsymbol{\mu}} = \frac{1}{n} \sum_{i=1}^{n} \mathbf{x}_i$$

**Anatomical** covariance matrix (estimated from cortical features)

# VNN vs PCA on age prediction task

Regression task

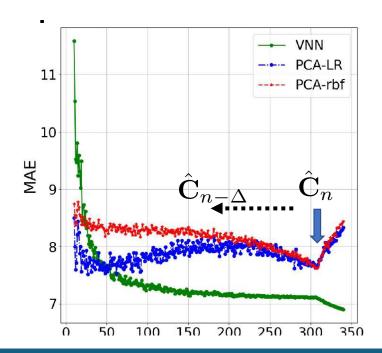


Cortical thickness — VNN — Estimate of age

Comparison against PCA-regression

**Data**: cortical thickness dataset (m = 104) from (n = 341) human subjects

➤ **Metric**: MAE (mean absolute error)

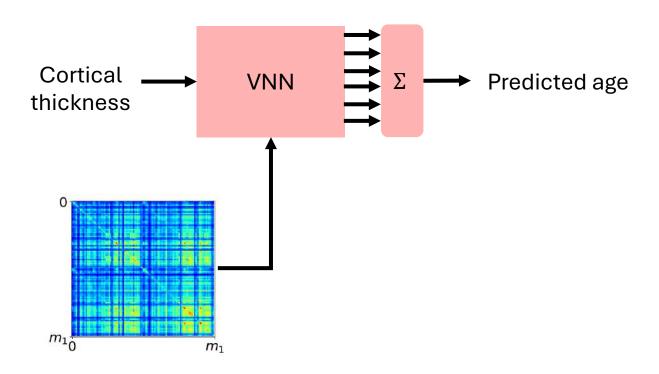


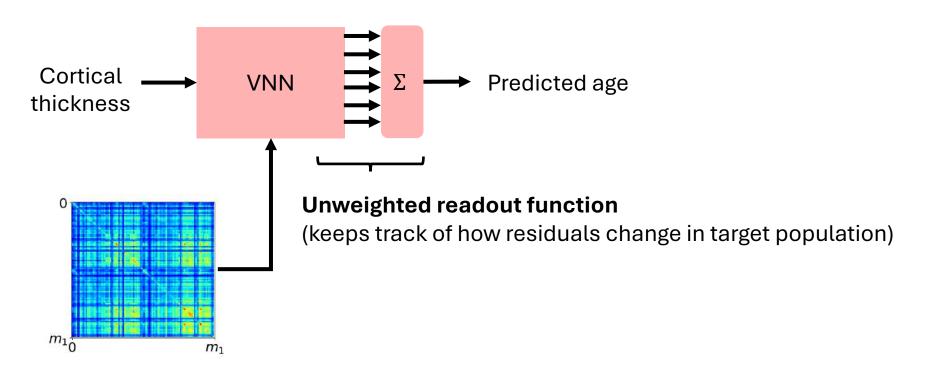
VNN: coVariance Neural Network

PCA-LR: PCA-regression with linear kernel

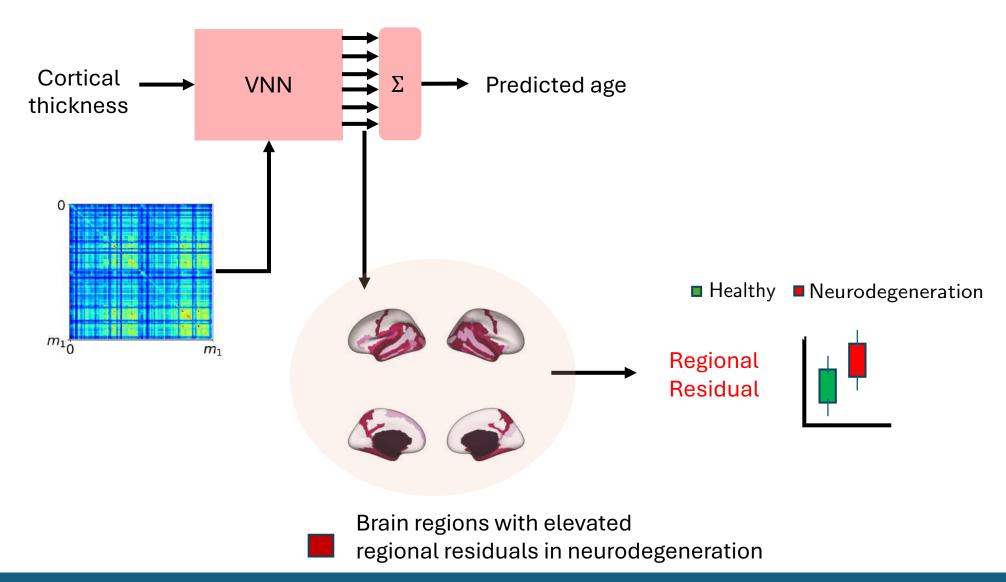
PCA-rbf: PCA regression with rbf kernel

VNN outperforms PCA and is **more stable** [Sihag et al., 2022]

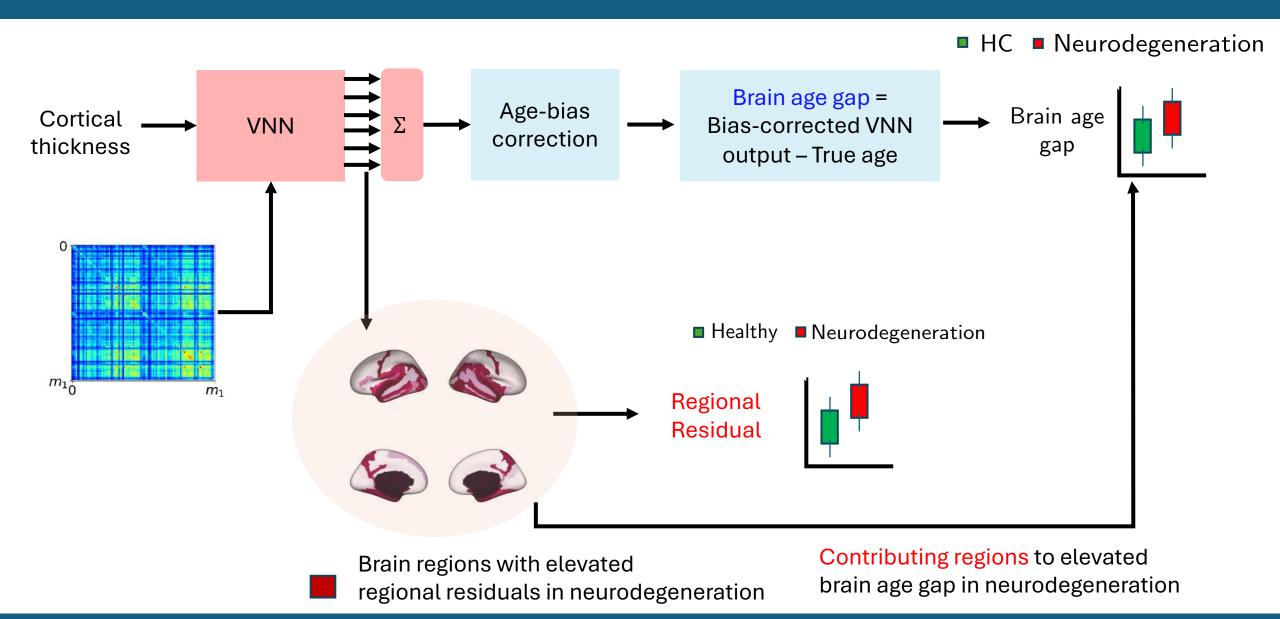




#### VNNs provide an anatomically interpretable and explainable brain age gap



#### VNNs provide an anatomically interpretable and explainable brain age gap



#### Experiments

Participants from OASIS-3 dataset, 148 cortical thickness features per individual

(Distrieux brain atlas)

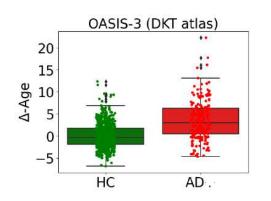
	HC	AD
Number	611	194
Age	68.38 (7.62)	74.72 (7.02)
Sex (m/f)	260/351	100/94
CDR sum of boxes	0	3.45 (1.74)

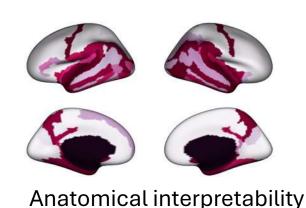
**HC group**: cognitively normal

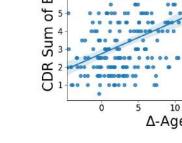
**AD group**: AD diagnosis

**CDR**: Clinical dementia rating

Brain age gap is elevated in AD group and correlated with CDR sum of boxes

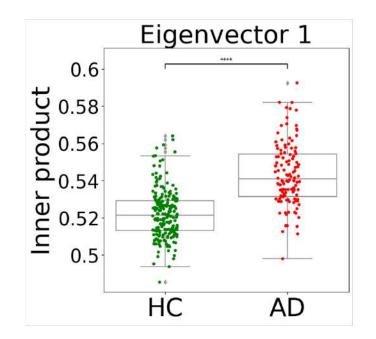


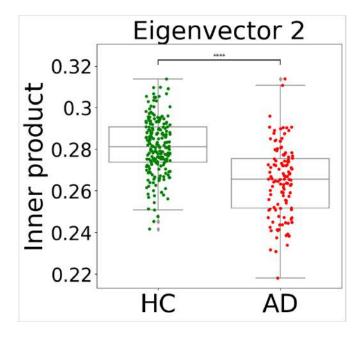




#### Experiments

VNN distinctly exploits eigenvectors in AD and HC groups



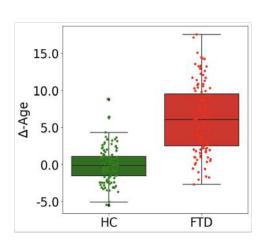


=> explains anatomical interpretability of brain age gap in AD

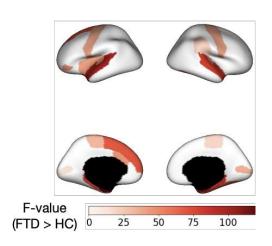
#### Experiments

- Whole brain cortical thickness dataset for Frontotemporal Dementia (FTD)
  - Healthy controls (HC, n = 114, age = 64.51 ± 6.51 years, 65 females)
  - FTD diagnosis (FTD, n = 119, age = 64.72 ± 6.78 years, 47 females)
- > 68 cortical thickness features (Desikan-Killiany atlas)

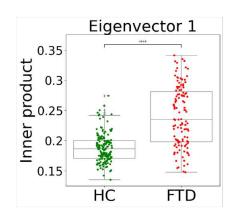
#### Brain age gap distributions

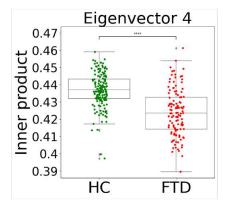


Anatomic interpretability

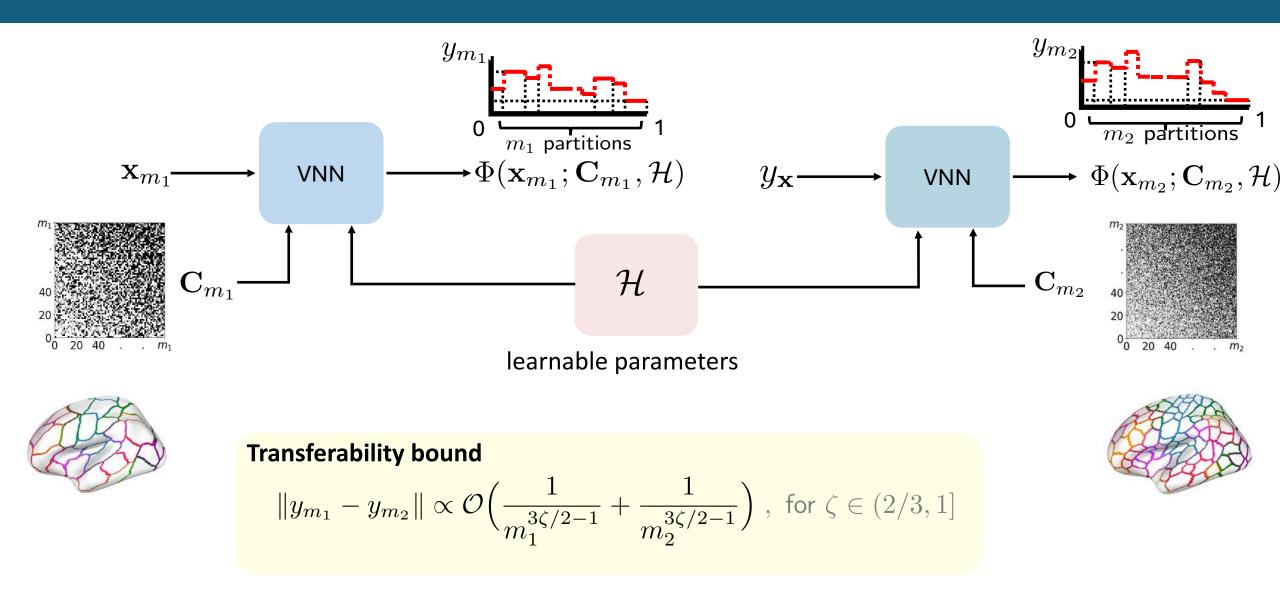


Explaining anatomic interpretability



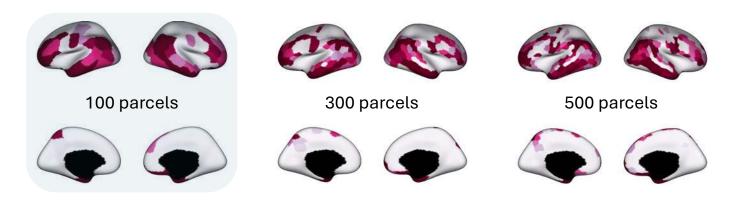


#### VNNs are provably transferable

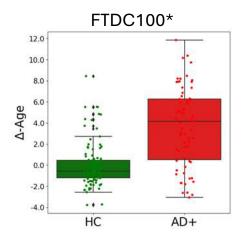


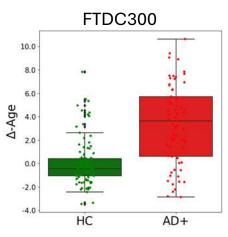
#### Recap: Transferability of VNNs cross-validates brain age gap in multi-resolution setting

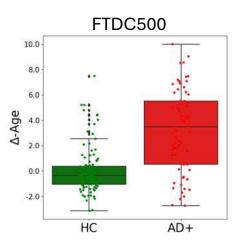
**Objective**: Brain age gap prediction in HC (healthy) and AD+ (Alzheimer's) cohorts from VNNs trained on 100-feature dataset



 ROIs contributing to elevated brain age gap in AD+ across different resolutions



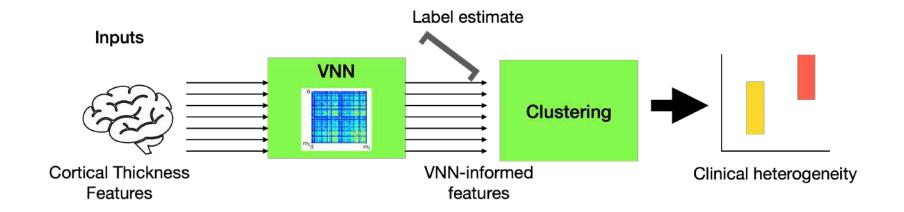




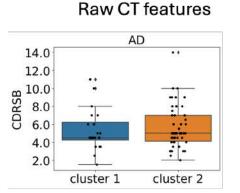
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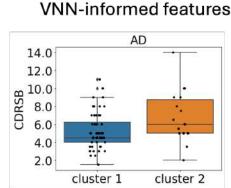
#### VNNs as pre-trained models...

Uncovering disease heterogeneity with VNNs as pre-trained models



VNNs offer more significant clinical stratification than raw anatomical features



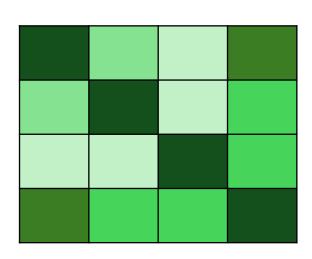


VNN enhances the clinical relevance of anatomical features

# Alignment of covariance with learning: An NTK perspective

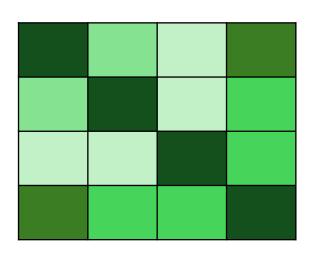
# Are covariance matrices suitable for a learning task?

- Covariance matrices add a meaningful inductive bias to neural nets
  - Covariance matrix captures the (linear) structure
- VNNs provide the bridge between PCA and GNNs



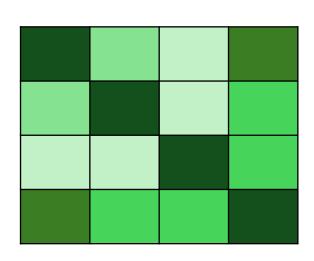
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  - Is performance good?
  - good generalization?



# Are covariance matrices suitable for a learning task?

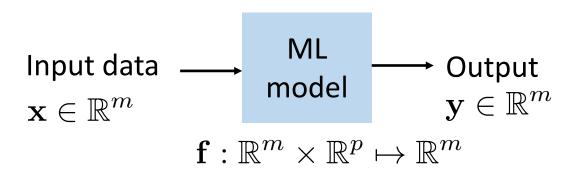
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- > VNNs provide the bridge between PCA and GNNs
- Can we quantify the suitability of covariance to learning objective?
  - Is performance good?
  - good generalization?
- > Neural tangent kernels (NTKs)-driven insights for VNNs



- NTKs describe the evolution of neural nets during training by gradient descent
- Example:

Predict y from x using

ML model f (parameters h)



- NTKs describe the evolution of neural nets during training by gradient descent
- Example:

Predict y from x using

ML model **f** (parameters **h**)

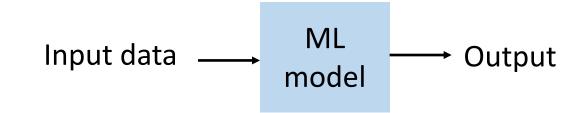
Input data 
$$\mathbf{x} \in \mathbb{R}^m$$
  $\longrightarrow$   $\mathbf{ML}$   $\mathbf{model}$   $\longrightarrow$   $\mathbf{Output}$   $\mathbf{y} \in \mathbb{R}^m$   $\mathbf{f} : \mathbb{R}^m \times \mathbb{R}^p \mapsto \mathbb{R}^m$ 

Loss:  $\mathcal{L} = \text{mean squared error (MSE)}$ 

$$= \sum_{i \in \mathsf{Data}} \|\mathbf{y}_i - \mathbf{f}(\mathbf{x}_i, \mathbf{h}))\|^2$$

Optimize parameters h using gradient descent

#### **Evolution of gradient descent**



(linear approximation)

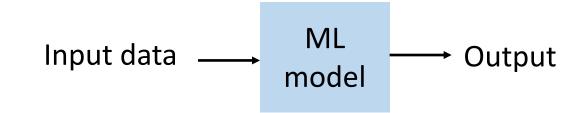
$$\mathbf{f}(\mathbf{x}, \mathbf{h}^{(t+1)}) = \mathbf{f}(\mathbf{x}, \mathbf{h}^{(t)}) - \eta \mathbf{\Theta}(\mathbf{x}, \mathbf{h}^{(t)}) \cdot (\mathbf{f}(\mathbf{x}, \mathbf{h}^{(t)}) - y)$$

$$\eta$$
: learning rate  $\Theta(\mathbf{x}, \mathbf{h}^{(t)})$ : NTK matrix

$$\Theta(\mathbf{x}_i, \mathbf{x}_j) = \nabla_{\mathbf{h}} \mathbf{f}(\mathbf{x}, \mathbf{h})^\mathsf{T} \nabla_{\mathbf{h}} \mathbf{f}(\mathbf{x}, \mathbf{h})$$

Depends on model architecture + input

Evolution of gradient descent



(linear approximation)

$$\mathbf{f}(\mathbf{x}, \mathbf{h}^{(t+1)}) = \mathbf{f}(\mathbf{x}, \mathbf{h}^{(t)}) - \eta \Theta(\mathbf{x}, \mathbf{h}^{(t)}) \cdot (\mathbf{f}(\mathbf{x}, \mathbf{h}^{(t)}) - y)$$

$$\eta : \text{ learning rate } \Theta(\mathbf{x}, \mathbf{h}^{(t)}) : \text{ NTK matrix } Depends on model$$

$$\Theta(\mathbf{x}_i, \mathbf{x}_i) = \nabla_{\mathbf{h}} \mathbf{f}(\mathbf{x}, \mathbf{h})^\mathsf{T} \nabla_{\mathbf{h}} \mathbf{f}(\mathbf{x}, \mathbf{h})$$
 architecture + input

 $\succ$  For neural networks with **infinite width**, NTK matrix is constant w.r.t  ${f h}$ 

$$\mathbf{\Theta}(\mathbf{x}, \mathbf{h}^{(t)}) \rightarrow \mathbf{\Theta}(\mathbf{x})$$

Convergence of gradient descent dictated by alignment between NTK and data

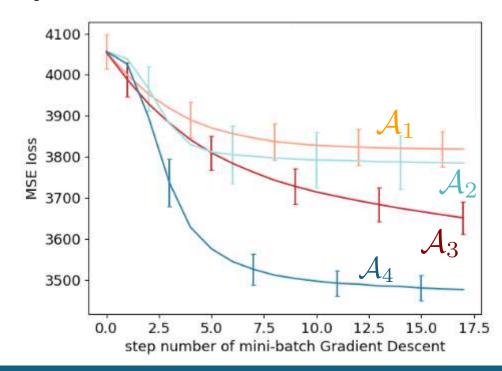
$$\|\mathbf{f}(\mathbf{x}, \mathbf{h}^{(t)}) - \mathbf{y}\|_2^2 \propto \mathcal{A} \text{ where } \mathcal{A} = \mathbf{y}^\mathsf{T} \mathbf{\Theta} \mathbf{y}$$
  
Larger  $\mathcal{A} \implies$  faster convergence

Convergence of gradient descent dictated by alignment between NTK and data

$$\|\mathbf{f}(\mathbf{x}, \mathbf{h}^{(t)}) - \mathbf{y}\|_2^2 \propto -\mathcal{A}$$
 where  $\mathcal{A} = \mathbf{y}^\mathsf{T} \mathbf{\Theta} \mathbf{y}$ 

Larger  $\mathcal{A}$   $\Longrightarrow$  faster convergence

#### **Example**

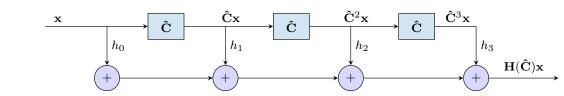


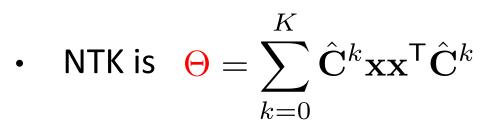
$$A_1 < A_2 < A_3 < A_4$$

Larger alignment implies better convergence/ loss

#### NTK for covariance filter

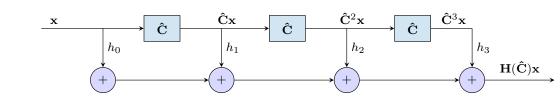
ightharpoonup For a covariance filter  $\mathbf{H}(\hat{\mathbf{C}}) = \sum h_k \hat{\mathbf{C}}^k$ ,





### Alignment for covariance filter

ightharpoonup For a covariance filter  $\mathbf{H}(\hat{\mathbf{C}}) = \sum h_k \hat{\mathbf{C}}^k$ ,



• NTK is 
$$\Theta = \sum_{k=0}^K \hat{\mathbf{C}}^k \mathbf{x} \mathbf{x}^\mathsf{T} \hat{\mathbf{C}}^k$$

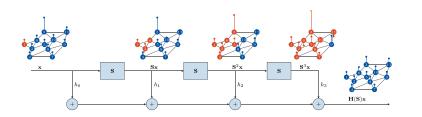
convergence of learning with covariance filter is dictated by

$$\mathcal{A} = \mathbf{y}^\mathsf{T} \left( \sum_{k=0}^K \hat{\mathbf{C}}^k \mathbf{x} \mathbf{x}^\mathsf{T} \hat{\mathbf{C}}^k \right) \mathbf{y}$$

Alignment between x,  $\hat{C}$ , and y

# Data-driven graph by optimizing alignment

- > Treating alignment as an optimization objective
  - Goal: find the optimal graph shift operator



$$\mathbf{S}^* = \max_{\mathbf{S}} \mathcal{A}(\mathbf{S})$$

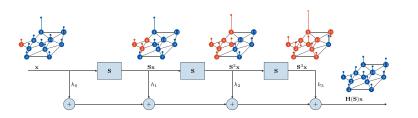
$$= \max_{\mathbf{S}} \mathbf{y}^\mathsf{T} \left( \sum_{k=0}^K \mathbf{S}^k \mathbf{x} \mathbf{x}^\mathsf{T} \mathbf{S}^k \right) \mathbf{y}$$

Find **S** that maximizes

$$\sum_{k=0}^{K} (\mathbf{y}^\mathsf{T} \mathbf{S} \mathbf{x})^2$$

### Data-driven graph by optimizing alignment

- Treating alignment as an optimization objective
  - **Goal**: find the *optimal* graph shift operator



$$\mathbf{S}^* = \max_{\mathbf{S}} \mathcal{A}(\mathbf{S})$$

$$= \max_{\mathbf{S}} \mathbf{y}^{\mathsf{T}} \left( \sum_{k=0}^{K} \mathbf{S}^k \mathbf{x} \mathbf{x}^{\mathsf{T}} \mathbf{S}^k \right) \mathbf{y}$$

Find **S** that maximizes

• Graph shift operator 
$$\mathbf{S}$$
• Input  $\mathbf{x}$ 
• Output  $\mathbf{v}$ 

Correlation between

- Output y

# Data-driven covariance graph by optimizing alignment

> Cross-covariance graph optimizes alignment

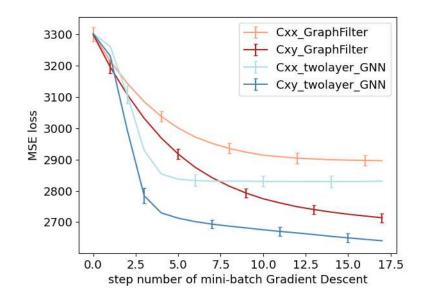
$$\mathbf{S}^* = \frac{1}{2}(\mathbf{x}\mathbf{y}^\mathsf{T} + \mathbf{y}\mathbf{x}^\mathsf{T})$$

# Data-driven covariance graph by optimizing alignment

> Cross-covariance graph optimizes alignment

$$\mathbf{S}^* = \frac{1}{2}(\mathbf{x}\mathbf{y}^\mathsf{T} + \mathbf{y}\mathbf{x}^\mathsf{T})$$

- Numerical results
  - Time series forecasting: predicting next time step



- Cross-covariance achieves better loss
- GNN with cross-covariance outperforms VNN

#### Variants of VNNs

#### Are VNNs enough?

- Limitations of VNNs
  - Sample covariance could be poor quality in low data, high dimensionality setting
  - High computational cost (quadratic in size for dense covariance)
  - No considerations of temporal, evolving data
  - Prone to undesired bias within the data

#### Low data, high dimensional settings

Sample covariance matrix is dense

 $\Longrightarrow$  **noisy** entries in low data, high dimensional settings

computationally inefficient VNNs (quadratic complexity)

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    - Improve estimation quality
    - Common in real world

(brain imaging, finance, etc.)

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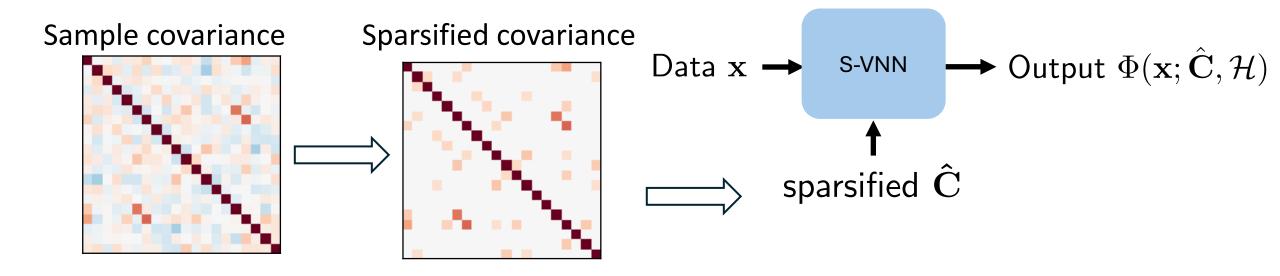
Improve computational efficiency

brain imaging, finance, etc.

For generic covariance:

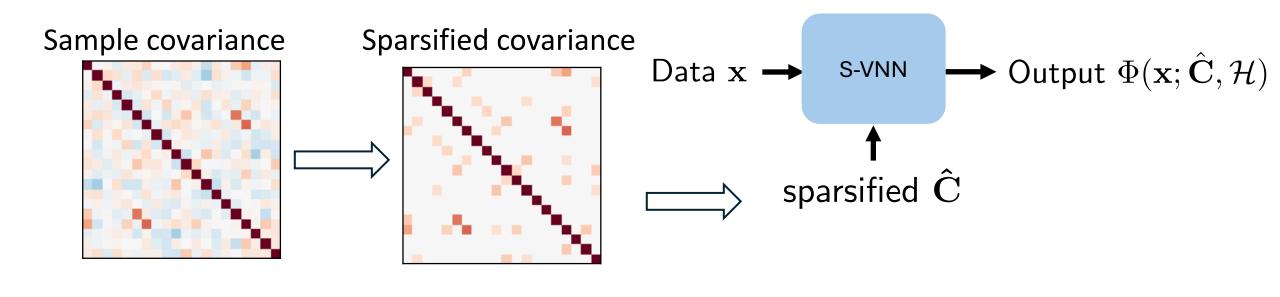
#### Sparse VNNs

> Sparse VNNs: sparsify the covariance matrix with thresholding techniques



#### Sparse VNNs

Sparse VNNs: sparsify the covariance matrix with thresholding techniques



What thresholding techniques?

Are sparse VNNs stable?

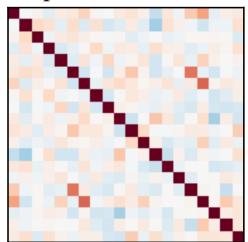
questions to address

### Hard thresholding

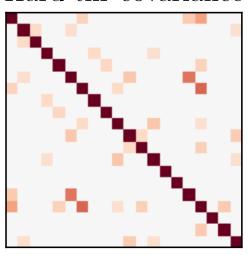
#### Definition

$$\eta(\hat{\mathbf{C}})_{ij} = \hat{c}_{ij} \text{ if } |\hat{c}_{ij}| \ge \tau/\sqrt{n}, 0 \text{ otherwise}$$

Empirical covariance



Hard-thr covariance



### Hard thresholding

#### Definition

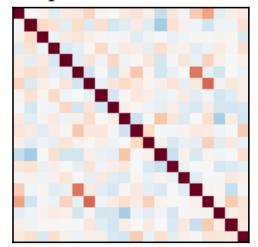
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Stability bound

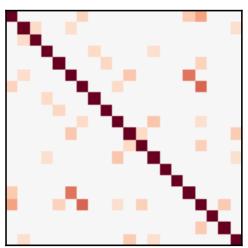
$$\|\mathbf{H}(\hat{\mathbf{C}}_{\mathsf{thr}}) - \mathbf{H}(\hat{\mathbf{C}})\| = \mathcal{O}\left(\frac{c_0}{n^{1/2}}\right)$$

 $c_0$ : number of non-zero elements in  $\hat{\mathbf{C}}_{\mathsf{thr}}$ 

Empirical covariance



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### Hard thresholding

#### Definition

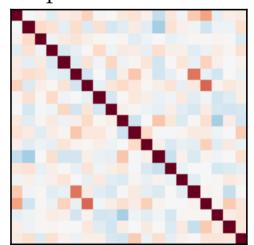
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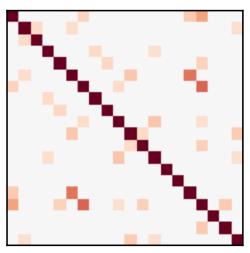
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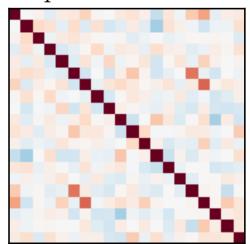
> Stability bound for S-VNNs is **tighter** than *dense* VNNs

### Soft thresholding

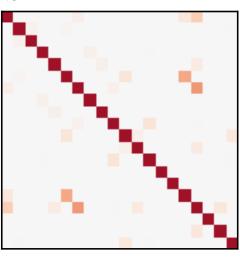
#### Definition

$$\eta(\hat{\mathbf{C}})_{ij} = \hat{c}_{ij} - \operatorname{sign}(\hat{c}_{ij})\tau/n \text{ if } |\hat{c}_{ij}| \geq \tau/\sqrt{n}, 0 \text{ otherwise}$$

Empirical covariance



Soft-thr covariance



#### Soft thresholding

#### Definition

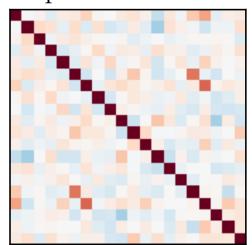
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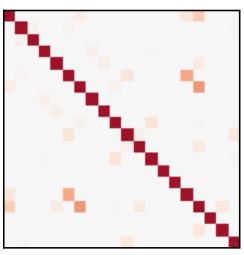
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 $c_0$ : number of non-zero elements in  $\hat{\mathbf{C}}_{\mathsf{thr}}$ 

Empirical covariance



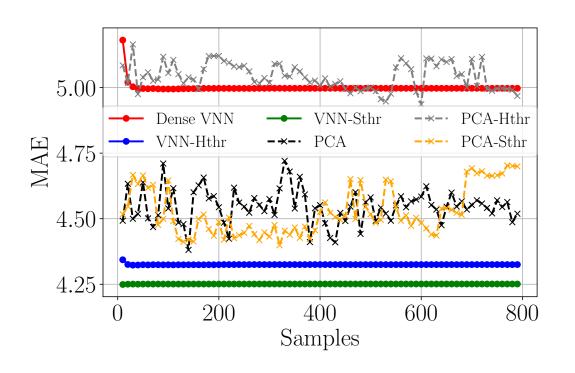
Soft-thr covariance



Stability bound for S-VNNs is tighter than dense VNNs

### Sparse VNNs: Numerical results

 Train VNNs/PCA on one covariance and test on another covariance estimated from less samples (synthetic dataset)

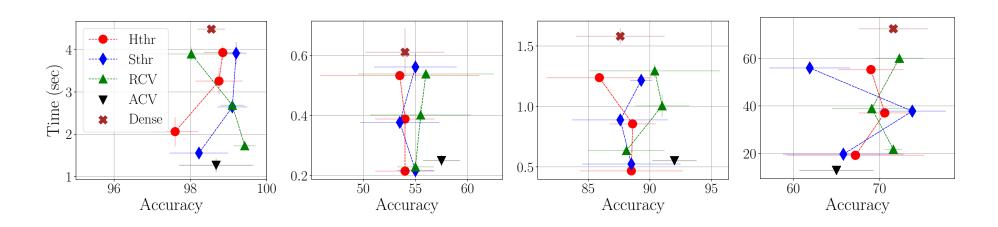


#### **Results**

- S-VNN (both soft and hard thresholding) **outperform** PCA and *dense* VNNs
- VNNs more stable than PCA

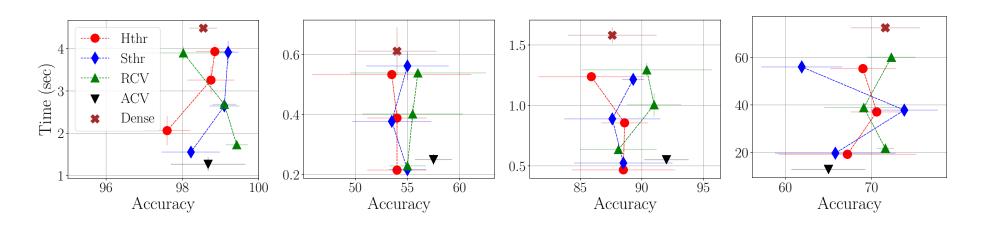
### Sparse VNNs: Numerical results

- Classification task on real data
- Datasets (from left to right)
  - Brain recordings: Epilepsy and CNI classify patient condition
  - Human action recognition: MHEALTH and Realdisp classify action



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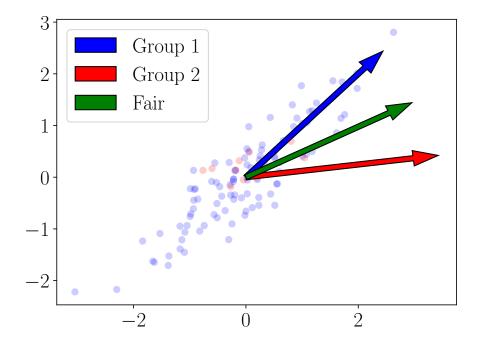
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S-VNNs are faster and achieve better performance than dense VNNs

### Limitations of VNNs - 2

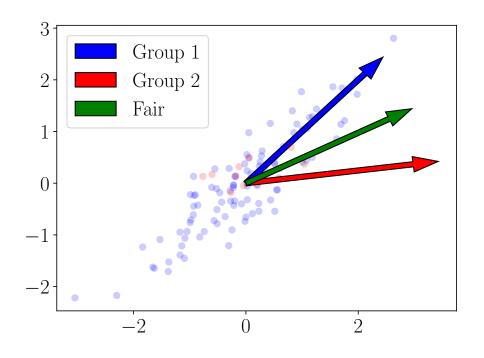
- Datasets may contain harmful biases
  - For e.g., under-represented groups
  - Biased (unfair) performance
  - Fair PCA might be unstable



### Limitations of VNNs - 2

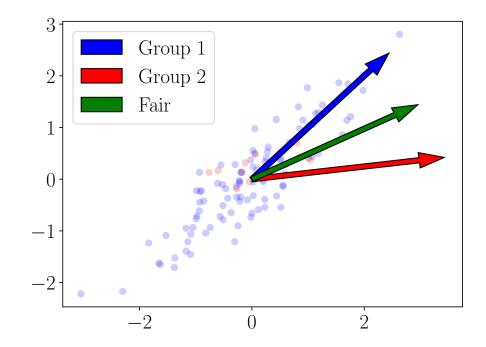
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- Fair VNNs (F-VNNs)



- Fairness: parity in performance across groups within data
- How to make VNNs fair?
- Are Fair VNNs stable?

questions to address

#### Fair covariance estimates

#### Balanced covariance

For two groups g and h,

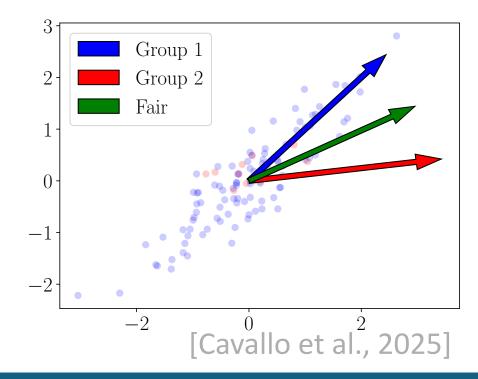
$$\hat{\mathbf{C}}_{\mathsf{bal}} = \alpha \hat{\mathbf{C}} + (1 - \alpha)(\hat{\mathbf{C}}_h - \hat{\mathbf{C}}_g) = \alpha_g \hat{\mathbf{C}}_g + \alpha_h \hat{\mathbf{C}}_h$$

#### Debiased covariance

$$\hat{\mathbf{C}}_{\mathsf{deb}} = \mathbf{X}^{\mathsf{T}} (\mathbf{I}_m + \beta \mathbf{Z} \mathbf{Z}^{\mathsf{T}})^{-1} \mathbf{X} / n$$

X: data matrix

**Z**: groups of samples



## Bias-mitigation penalty

> F-VNNs are trained with a **loss penalty** that encourages fairness

$$\min_{\mathcal{H}} \ \gamma \mathcal{L}(\mathbf{X}, \mathbf{y}, \Phi) + (1 - \gamma) \mathcal{R}(\mathbf{X}, \mathbf{y}, \mathbf{z}, \Phi)$$

 $\mathcal{L}$ : task-specific loss (for e.g., cross-entropy, MAE)

 $\mathcal{R}$ : bias penality (for e.g., performance difference across groups)

 $\gamma$ : balancing term

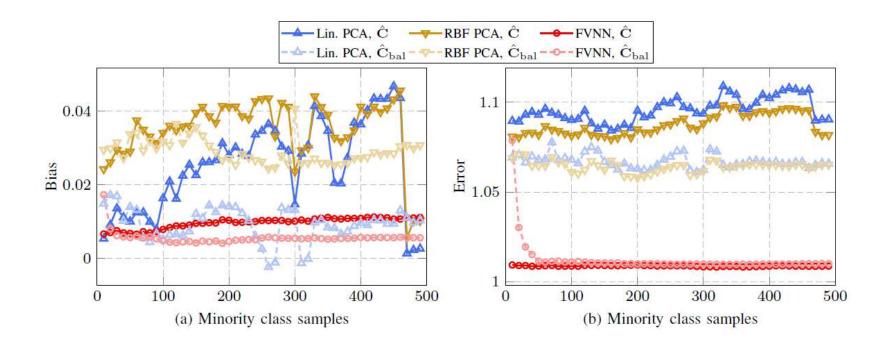
# Stability of F-VNNs

- Fair covariance estimates
  - $\hat{\mathbf{C}}_{\mathsf{deb}}$  and  $\hat{\mathbf{C}}_{\mathsf{bal}}$  are subject to covariance estimation errors
  - PCA with fair covariance estimates (Fair PCA) may be unstable
    - biased treatment

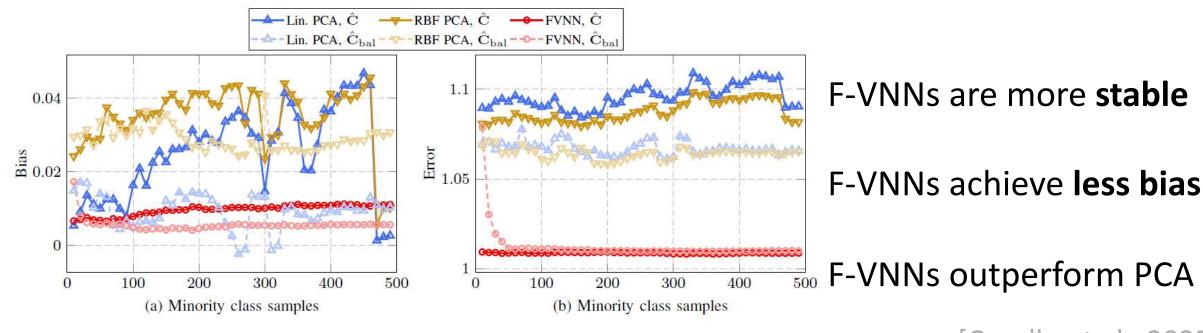
# Stability of F-VNNs

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    - biased treatment
- F-VNNs are stable
  - Stability of F-VNNs with **balanced** covariance  $\propto \mathcal{O}\left(\frac{1}{n_g^{1/2}}\right) + \mathcal{O}\left(\frac{1}{n_h^{1/2}}\right)$
  - Stability of F-VNNs with **debiased** covariance  $\propto \mathcal{O}\left(\frac{1}{n^{1/2}}\right)$  [Cavallo et al., 2025]

- Stability: synthetic biased data
  - Train on unbiased dataset
  - During test, replace covariance with unbalanced/fair version
  - Compare PCA+SVM with VNNs

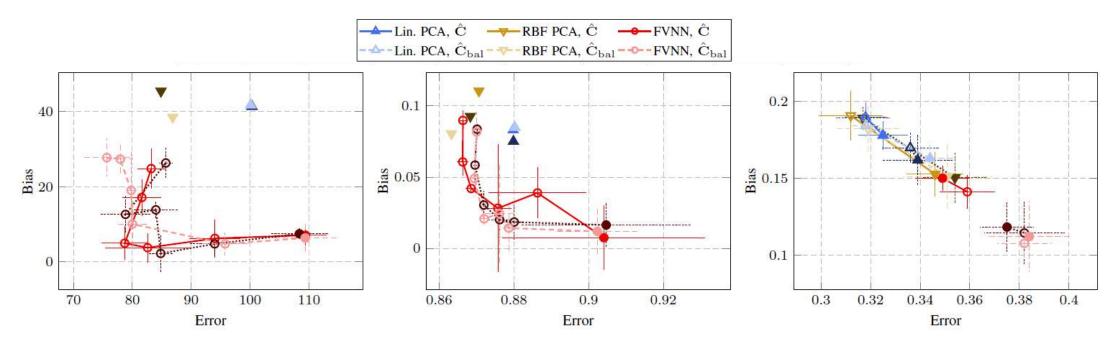


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#### Real world datasets

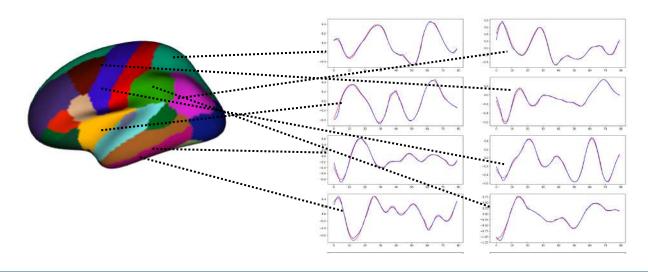
Dataset	Description	Task	Sensitive attribute
Parkinson (left)	Medical records of patients	Regression for Parkinson's level	Sex of patient
LSAC (center)	Law school students' features	Regression for GPA	Race of students
German credit (right)	Features of individuals applying for credit	Classification (good or bad)	Sex of individual



F-VNNs achieve better fairness and performance than PCA

### Limitations-3

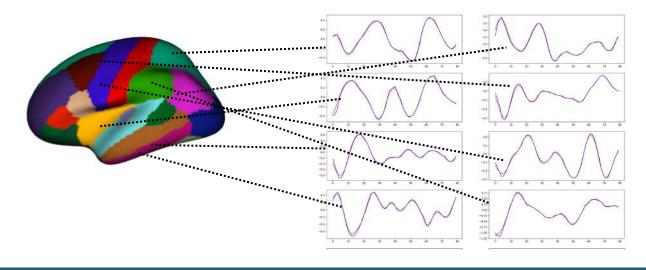
- > VNN models discussed so far operate on *static* data
  - Real world applications have dynamic data
  - Non-trivial modifications needed to handle temporal, non-stationary data
  - Online estimates introduce additional source of errors



### Limitations-3

- > VNN models discussed so far operate on *static* data
  - Real world applications have dynamic data
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  - Online estimates introduce additional source of errors

- Spatio-temporal VNNs (STVNNs)
  - VNNs for spatio-temporal datasets



#### Model design

Online covariance matrix estimate

$$\hat{\mathbf{C}}_{t+1} = \zeta_t \hat{\mathbf{C}}_t + \beta_t (\mathbf{x}_{t+1}) (\mathbf{x}_{t+1})^\mathsf{T}$$

#### Model design

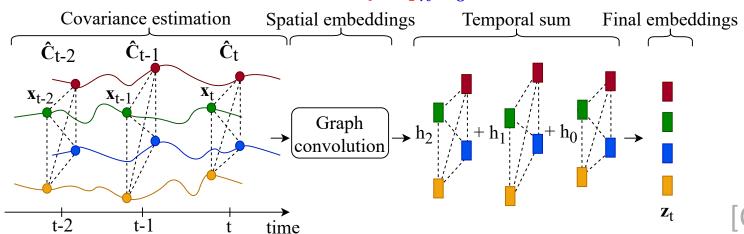
Online covariance matrix estimate

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Spatio-temporal coVariance filter

$$\mathbf{z}_t := \mathbf{H}(\mathbf{\hat{C}}_t, \mathbf{h}_t, \mathbf{x}_{T:t}) = \sum_{t'=0}^{T-1} \sum_{k=0}^{K} h_{kt'} \mathbf{\hat{C}}_t^k \mathbf{x}_{t-t'}$$

Spatial and temporal convolution



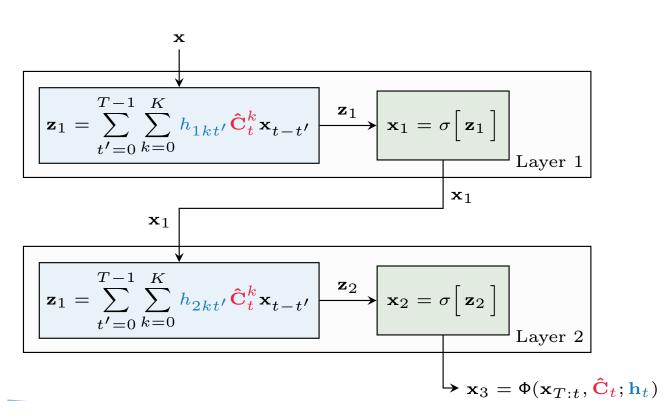
#### STVNN

Sequences of spatio-temporal covariance filters followed by non-linearity

$$\mathbf{z}_t^l = \sigma\left(\mathbf{H}^l(\mathbf{\hat{C}}_t, \mathbf{h}_t, \mathbf{z}_{T:t}^{l-1})\right)$$

Online parameter updates

$$\mathbf{h}_{t+1} = \mathbf{h}_t - \eta \nabla_t \mathcal{L}(\Phi(\mathbf{x}_{T:t}, \hat{\mathbf{C}}_t; \mathbf{h}_t))$$



#### STVNN

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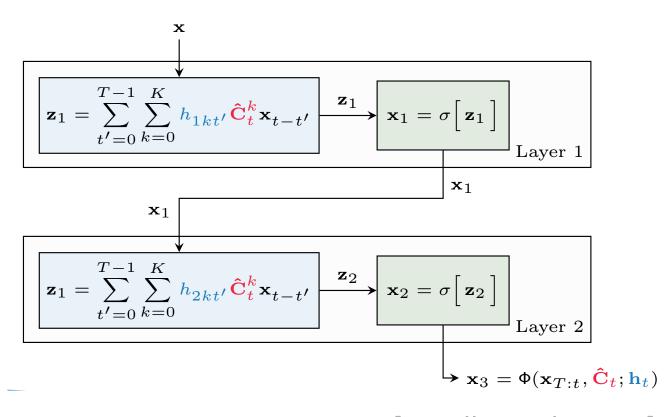
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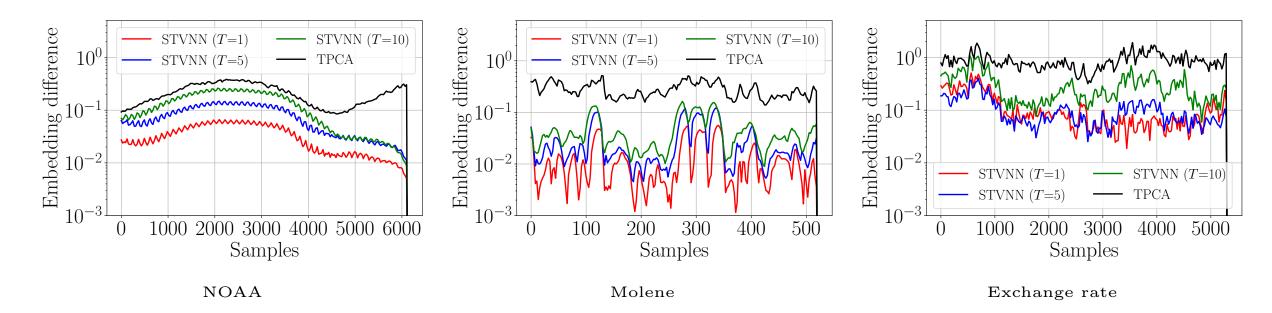
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STVNNs are stable

Stability bound 
$$\propto \mathcal{O}\left(\frac{1}{\sqrt{n}}\right)$$

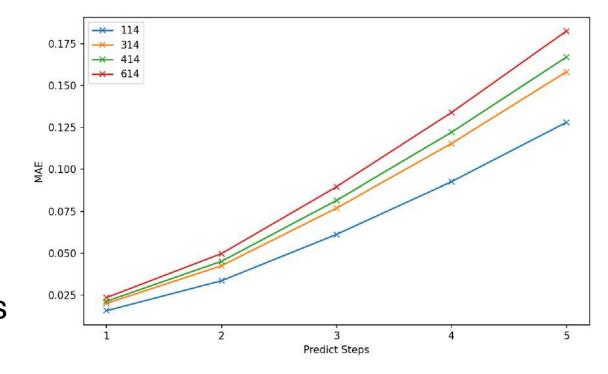


- Time series forecasting task (weather data and currency exchange rates)
  - Train with one covariance, test with another estimated from fewer samples

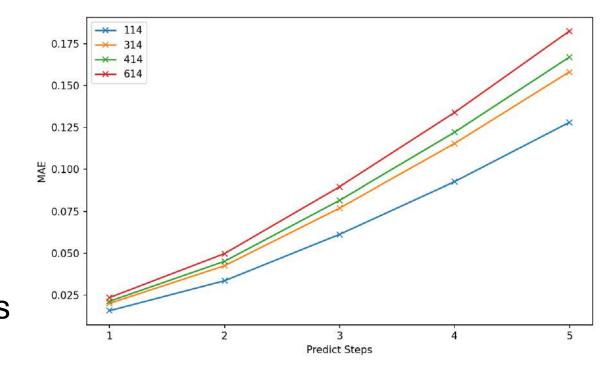


STVNNs are more stable than temporal-PCA (TPCA) Higher T (temporal window size), lower stability

- > Time series forecasting task (brain imaging data)
  - Data: HCP Young-adult dataset
  - BOLD data at at spatial scales
     of 114, 314, 414, 614 (Schaefer's)
  - Train model on 314 resolution
    - Test on 114, 414, 614 resolutions



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STVNN demonstrates transferability across multi-scale spatio-temporal datasets

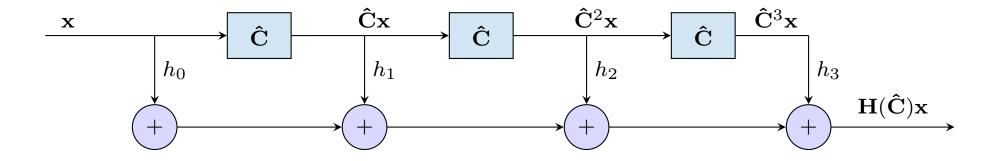
#### Conclusions and Future Directions

### Covariance filters

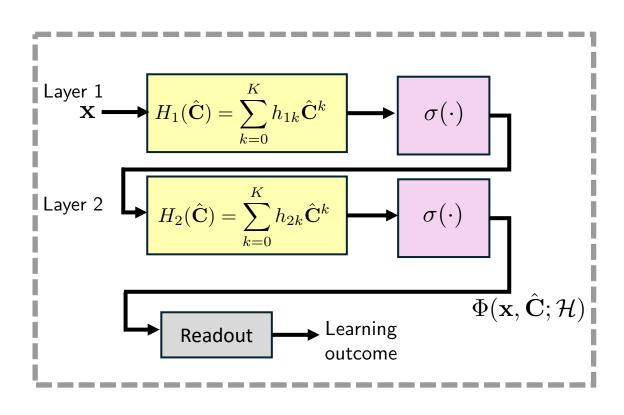
ightarrow A covariance filter is a polynomial in the covariance matrix  $\hat{\mathbf{C}}$ 

$$\mathbf{H}(\hat{\mathbf{C}}) = \sum_{k=0}^{K} h_k \hat{\mathbf{C}}^k \mathbf{x}$$

 $\triangleright$  We train the filter coefficients  $h_k$  to accomplish some task

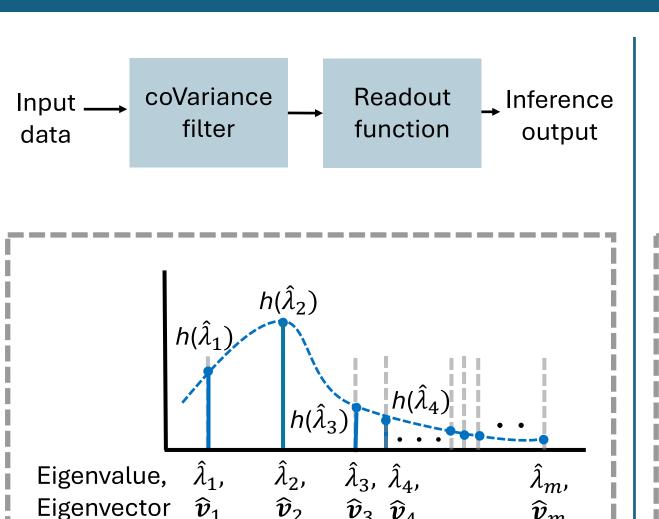


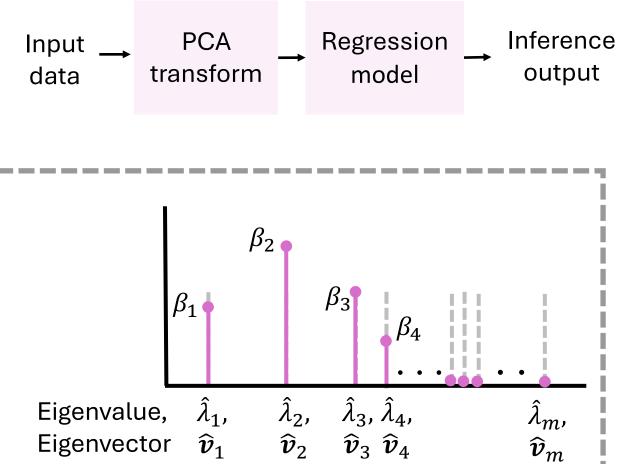
### CoVariance Neural Networks (VNNs)



- A VNN is a composition of layers
- > Each of which is a composition of
  - ... a covariance filter
  - ... with a pointwise nonlinearity
- $\blacktriangleright \Phi(\mathbf{x}; \hat{\mathbf{C}}, \mathcal{H})$  represents VNN output
- $\succ \mathcal{H}$  is the set of trainable filter taps

## Covariance Filters are Implicitly Equivalent to PCA



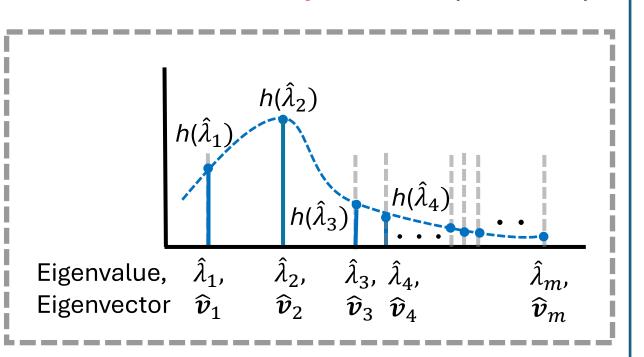


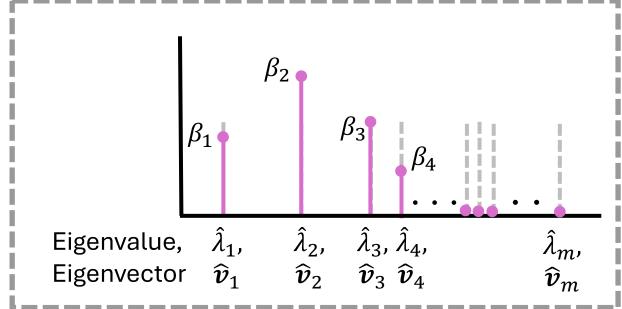
### Covariance Filters are Implicitly Equivalent to PCA

The difference is that covariance filters (and VNNs) do not require eigenvectors

Stability: Leading to more stable signal processing

Transferability: And the possibility of transferring trained filters across scales





### Stability of coVariance filters and Neural Networks

Outputs of coVariance filters NNs on true and estimated covariances are close

$$\left\| \mathbf{H}(\hat{\mathbf{C}}) - \mathbf{H}(\mathbf{C}) \right\| = \mathcal{O}\left(\frac{1}{n^{1/2 - \varepsilon}}\right) = \alpha_n \qquad \left\| \Phi(\mathbf{x}, \hat{\mathbf{C}}; \mathcal{H}) - \Phi(\mathbf{x}, \mathbf{C}; \mathcal{H}) \right\| \le LF^{L-1}\alpha_n$$

Provided that the filters (at each layer) have Lipschitz frequency responses

$$|h(\lambda_i) - h(\lambda_j)| \le Q \frac{|\lambda_i - \lambda_j|}{k_i}$$

> This requirement limits discriminability but it is a necessary limit

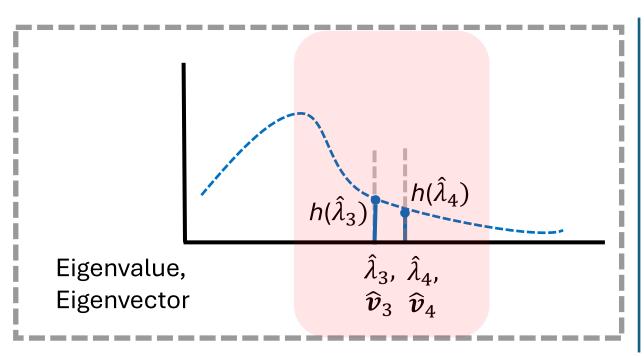
[Sihag et al., 2022]

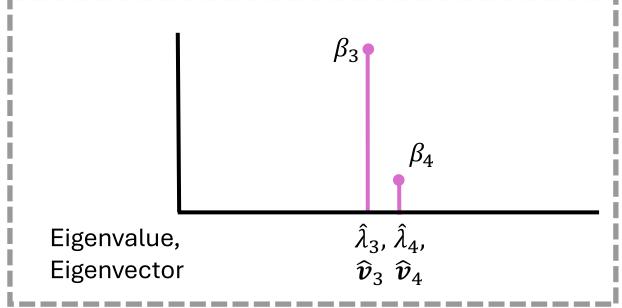
### PCA responds catastrophically to eigenspace estimation

Difference between true and estimated eigenvectors can be arbitrarily different

$$\|\hat{\mathbf{V}}\mathbf{x} - \mathbf{V}\mathbf{x}\| = \mathcal{O}\left(\frac{1}{n^{1/2}\min_{i \neq j} |\lambda_i - \lambda_j|}\right)$$

> Filters process similar eigenvalues with similar coefficients



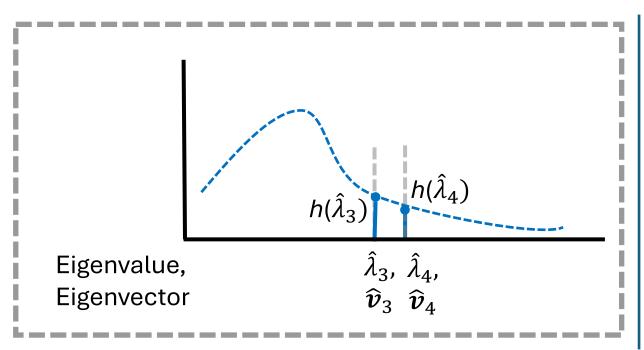


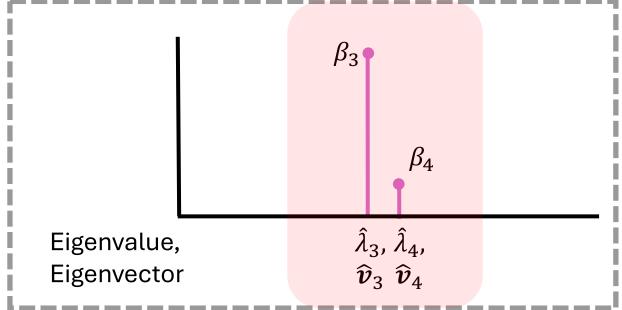
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> Eigenvectors with similar eigenvalues can be processed with dissimilar coefficients



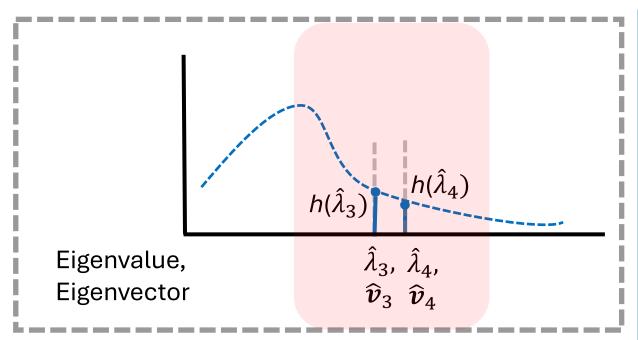


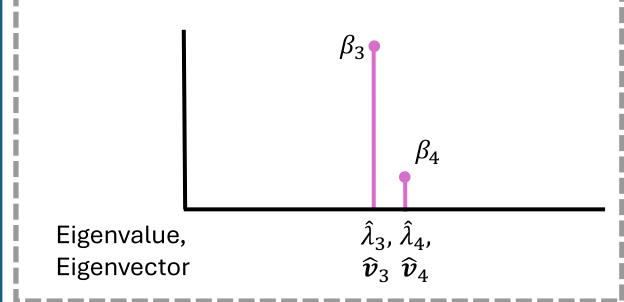
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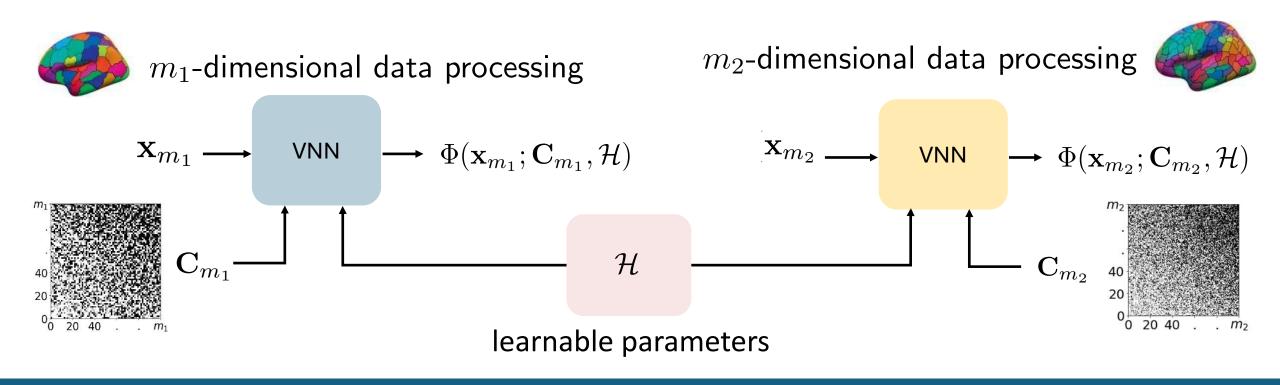
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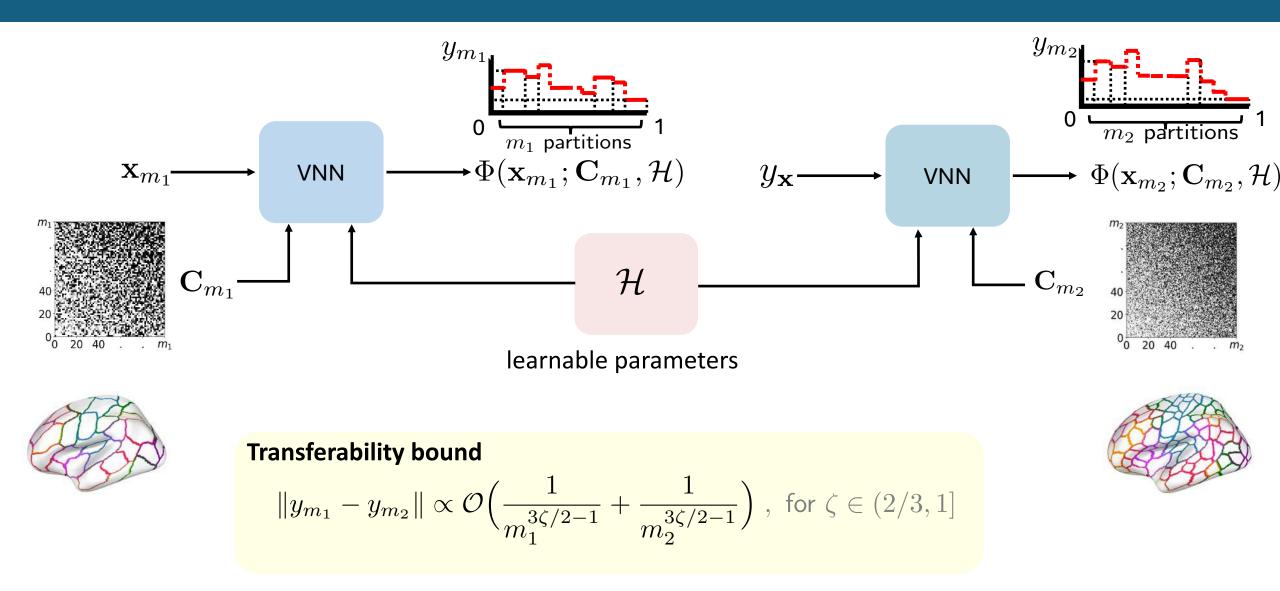


### coVariance Filters and VNNs are Scale-Free Models

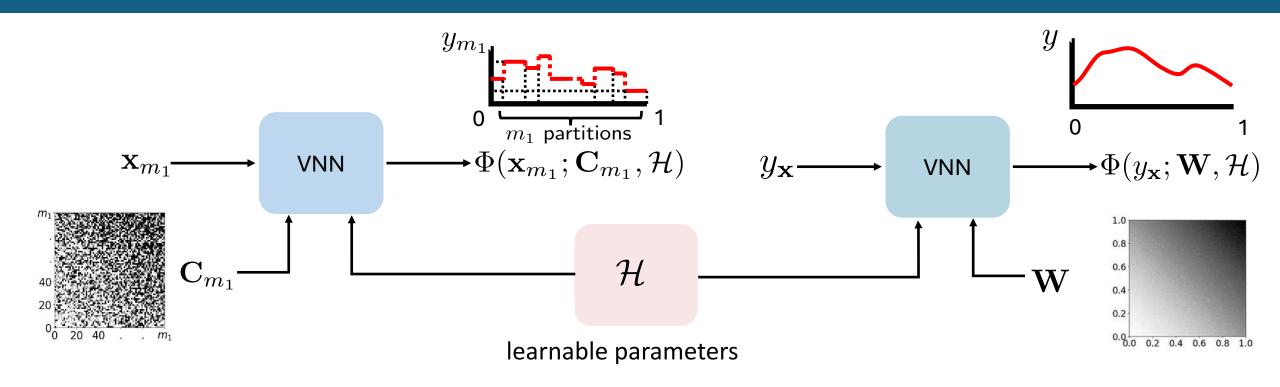
- > Filters and VNNs are defined by coefficients that we can transfer across scales
  - > Train at small scale and transfer to large scale
  - > Train jointly across a heterogeneous range of scales



### VNNs are provably transferable



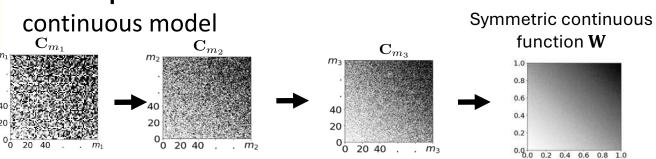
### VNNs are provably transferable to limit models





$$\|y_{m_1}-y\|\propto \mathcal{O}\left(rac{1}{m_1^{3\zeta/2-1}}
ight) \ , \ ext{for } \zeta\in (2/3,1]$$

\*Assumption: data is a discretization of a common



# VNNs are well suited for neuroimaging data analysis

> Properties of VNNs make them appealing for neuroimaging data analysis

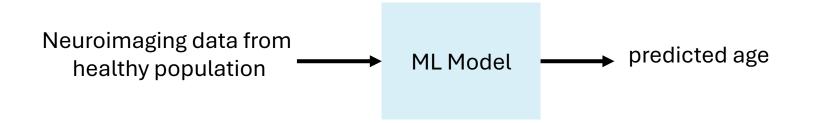
- Connections with PCA transparent outcomes by leveraging spectrum of covariance matrix
- Stability 
   reproducible outcomes in limited data settings

Transferability 
 — enhanced generalizability and robustness to choice of brain atlases

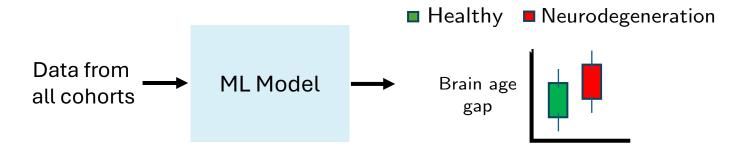
# Brain age gap prediction is a transfer learning problem

Train ML model to predict age on a large dataset (healthy population)

**Pre-training** 



Apply the pre-trained ML model on a target dataset (neurodegeneration)

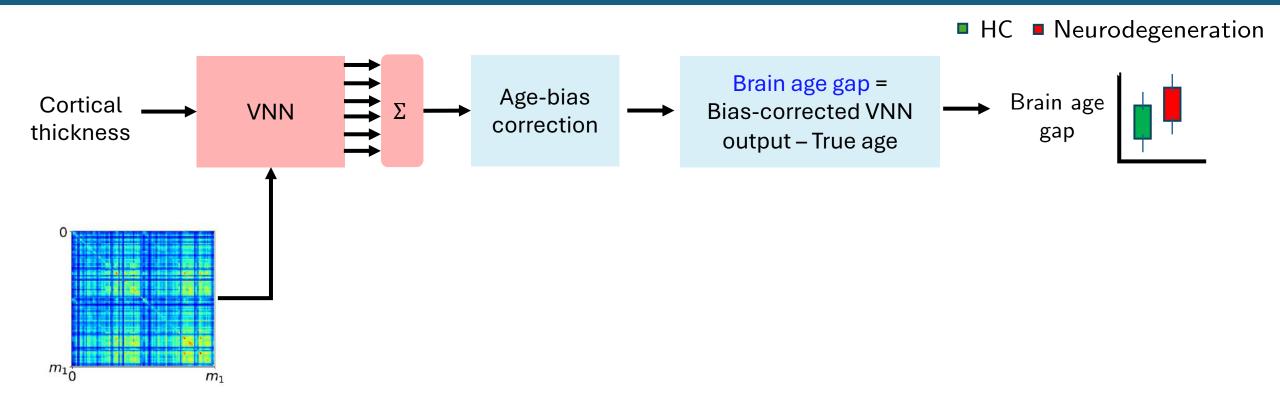


Brain age gap is the residual of the model

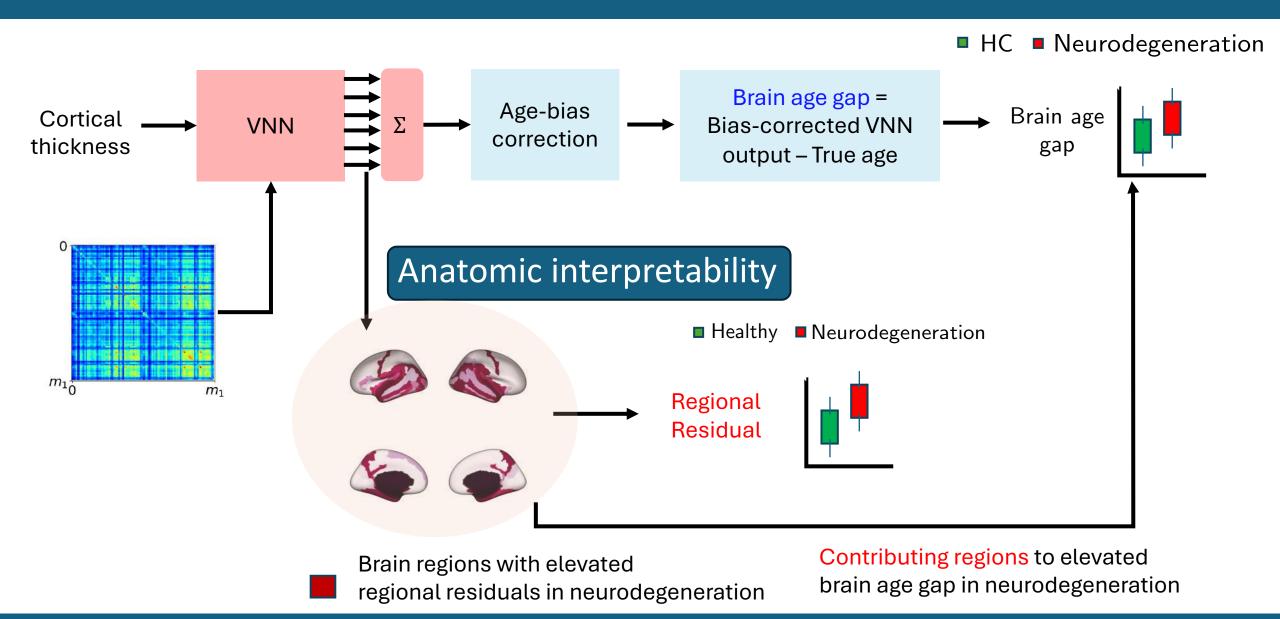
## A principled approach to brain age gap prediction

- > Focus on residuals of the ML model, not prediction performance
- > Qualitative evaluation during pre-training
  - what does the model learn during pre-training on healthy population?
- > Interpretability/explainability:
  - what's driving elevated brain age gap (residuals) in neurodegeneration?
- > Generalizability to diverse target populations

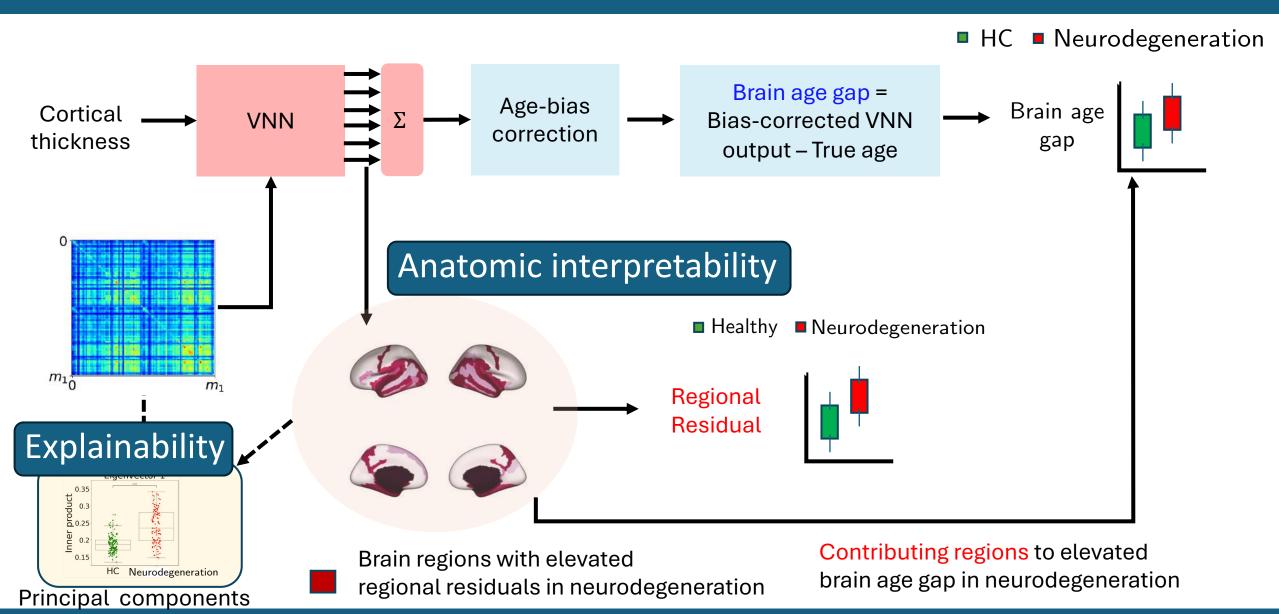
#### VNNs provide an anatomically interpretable and explainable brain age gap



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## **Concluding Remarks**

- Emerging areas
  - Sparse VNNs: sparsifying covariance matrix [Cavallo et al., 2024]
  - Spatiotemporal VNNs: temporal datasets [Cavallo et al., 2024]
  - Fair VNNs: unbiased outcomes with VNNs [Cavallo et al., 2025]
  - Optimality of covariance matrices: suitability of covariance to learning task [Khalafi et al., 2024]
  - Application to brain age gap prediction [Sihag et al., 2024; 2025]

### Future directions

- > Learning with **cross-covariance** graphs
  - Links with partial least squares/ canonical correlation analysis
- Expand interpretability/explainability of VNNs
  - How are eigenvectors exploited in STVNNs on dynamic datasets?
- Building interpretable biomarkers
  - Using other modalities (for e.g., fMRI)

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### Slides available at

