

Covariate Assisted Principal (**CAP**) Regression for Matrix Outcomes

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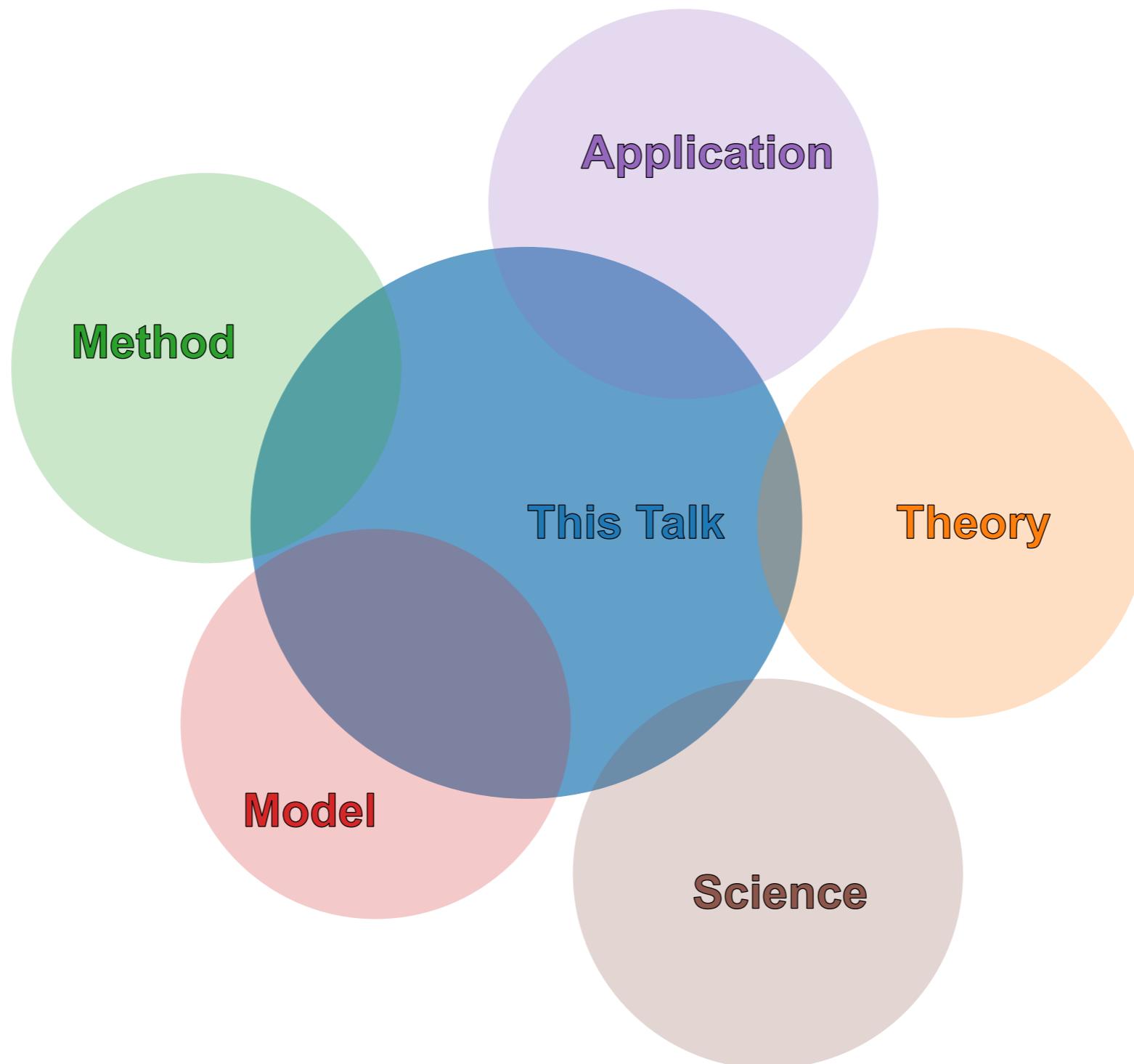


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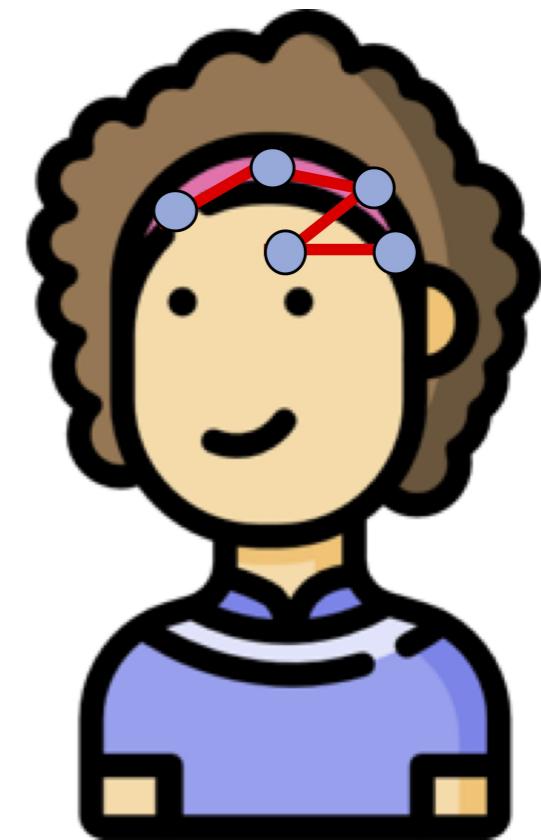
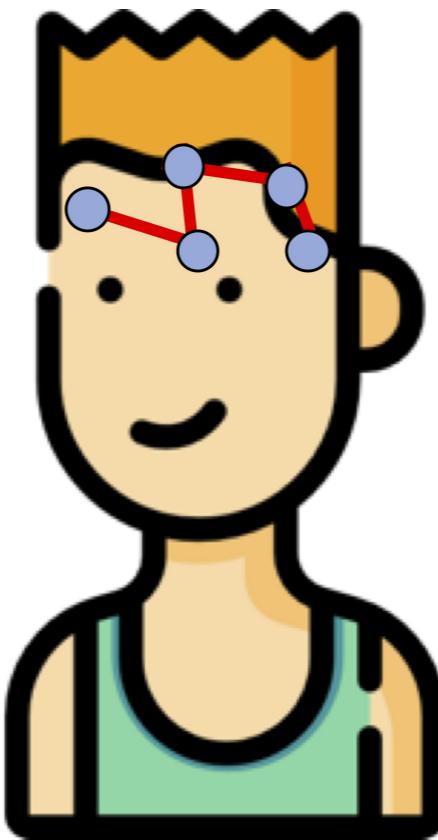
Slides viewable on web:
bit.ly/icsahz19



Statistics/Data Science Focuses



Motivating Example



Brain network connections vary by covariates (e.g. age/sex)

Goal: model how covariates change network connections

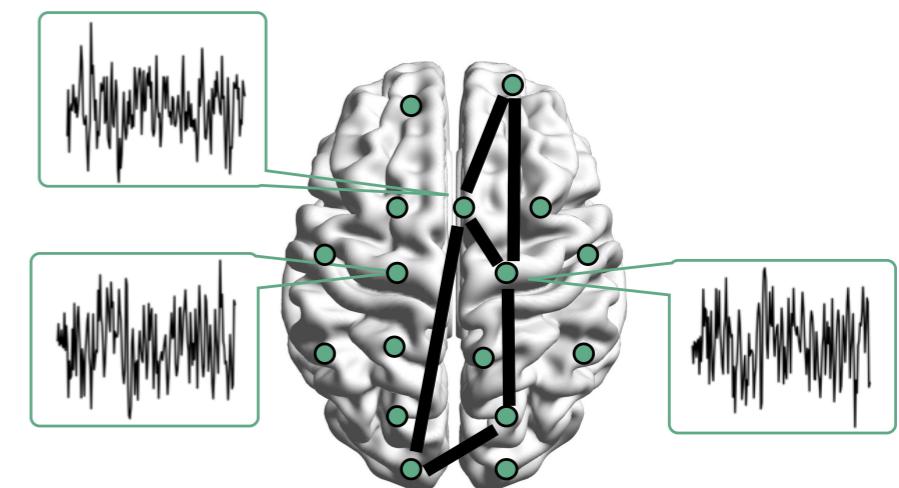
function(graph) = age $\times \beta_1 + \text{sex} \times \beta_2 + \dots$



Resting-state fMRI Networks



- fMRI measures brain activities over time
- Resting-state: "do nothing" during scanning
- Brain networks constructed using **cov/cor** matrices of time series



Mathematical Problem

- Given n (semi-)positive matrix outcomes,
 $\Sigma_i \in \mathbb{R}^{p \times p}$
- Given n corresponding vector covariates, $x_i \in \mathbb{R}^q$
- Find function $g(\Sigma_i) = x_i\beta, i = 1, \dots, n$
- In essence, **regress positive matrices on vectors**



Some Related Problems

- Heterogeneous regression or weighted LS:
 - Usually for scalar variance σ_i , find $g(\sigma_i) = f(x_i)$
 - Goal: to improve efficiency, not to interpret $x_i \beta$
- Covariance models [Anderson, 73; Pourahmadi, 99; Hoff, Niu, 12; Fox, Dunson, 15; Zou, 17]
 - Model $\Sigma_i = g(x_i)$, sometimes $n = i = 1$
 - Goal: better models for Σ_i
- Multi-group PCA [Flury, 84, 88; Boik 02; Hoff 09; Franks, Hoff, 16]
 - No regression model, cannot handle vector x_i
 - Goal: find common/uncommon parts of multiple Σ_i
- Tensor-on-scalar regression [Li, Zhang, 17; Sun, Li, 17]
 - No guarantees for positive matrix outcomes



Massive Edgewise Regressions

- Intuitive method by mostly neuroscientists
- Try $g_{j,k}(\Sigma_i) = \Sigma_i[j, k] = x_i \beta$
- Repeat for all $(j, k) \in \{1, \dots, p\}^2$ pairs
- Essentially $O(p^2)$ regressions for each connection
- Limitations: multiple testing $O(p^2)$, failure to account for dependencies between regressions



Our CAP in a Nutshell

$$\text{PCA}(\Sigma_i) = x_i \beta$$

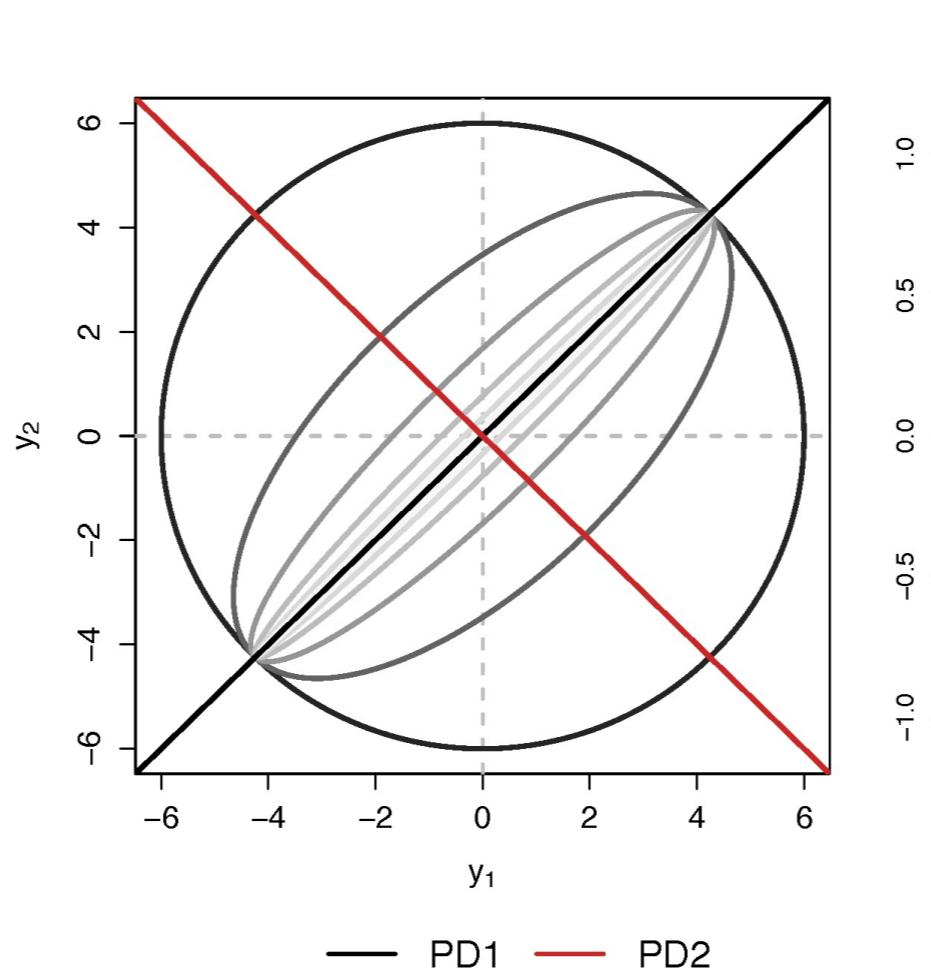
- Essentially, we aim to turn unsupervised PCA to a supervised PCA
- Ours differs from existing PCA methods:
 - Supervised PCA [Bair et al, 06] models **scalar-on-vector**



Model and Method

Model

- Find principal direction (PD) $\gamma \in \mathbb{R}^p$, such that:
$$\log(\gamma^\top \Sigma_i \gamma) = \beta_0 + x_i^\top \beta_1, \quad i = 1, \dots, n$$



Example ($p=2$): PD1 largest variation but not related to x
≡
PCA selects PD1, Ours selects **PD2**

Advantages

- Scalability: potentially for $p \sim 10^6$ or larger
- Interpretation: covariate assisted PCA
 - Turn **unsupervised** PCA into **supervised**
- Sensitivity: target those covariate-related variations
 - **Covariate assisted** SVD?
- Applicability: other big data problems besides fMRI



Method

- MLE with constraints:

$$\underset{\beta, \gamma}{\text{minimize}} \quad \ell(\beta, \gamma) := \frac{1}{2} \sum_{i=1}^n (x_i^\top \beta) \cdot T_i + \frac{1}{2} \sum_{i=1}^n \gamma^\top \Sigma_i \gamma \cdot \exp(-x_i^\top \beta),$$

such that $\gamma^\top H \gamma = 1$

- Two obvious constraints:

- C1: $H = I$
- C2: $H = n^{-1}(\Sigma_1 + \dots + \Sigma_n)$



Choice of H

Proposition: When (C1) $H = I$ in the optimization problem, for any fixed β , the solution of γ is the eigenvector corresponding to the minimum eigenvalue of matrix

$$\sum_{i=1}^n \frac{\Sigma_i}{\exp(x_i^\top \beta)}$$

Will focus on the constraint (C2)



Algoirthm

- Iteratively update β and then γ
- Prove explicit updates
- Extension to multiple γ :
 - After finding $\gamma^{(1)}$, we will update Σ_i by removing its effect
 - Search for the next PD $\gamma^{(k)}, k = 2, \dots$
 - Impose the orthogonal constraints such that γ^k is orthogonal to all $\gamma^{(t)}$ for $t < k$



Theory for β

Theorem: Assume $\sum_{i=1}^n x_i x_i^\top / n \rightarrow Q$ as $n \rightarrow \infty$. Let $T = \min_i T_i$, $M_n = \sum_{i=1}^n T_i$, under the true γ , we have

$$\sqrt{M_n} (\hat{\beta} - \beta) \xrightarrow{\mathcal{D}} \mathcal{N}(0, 2Q^{-1}), \quad \text{as } n, T \rightarrow \infty,$$

where $\hat{\beta}$ is the maximum likelihood estimator when the true γ is known.

Theory for γ

Theorem: Assume $\Sigma_i = \Gamma \Lambda_i \Gamma^\top$, where $\Gamma = (\gamma_1, \dots, \gamma_p)$ is an orthogonal matrix and $\Lambda_i = \text{diag}\{\lambda_{i1}, \dots, \lambda_{ip}\}$ with $\lambda_{ik} \neq \lambda_{il}$ ($k \neq l$), for at least one $i \in \{1, \dots, n\}$. There exists $k \in \{1, \dots, p\}$ such that for $\forall i \in \{1, \dots, n\}$ $\gamma_k^\top \Sigma_i \gamma_k = \exp(x_i^\top \beta)$. Let $\hat{\gamma}$ be the maximum likelihood estimator of γ_k in Flury, 84. Then assuming that the assumptions are satisfied, $\hat{\beta}$ from our algorithm is $\sqrt{M_n}$ -consistent estimator of β .



Simulations



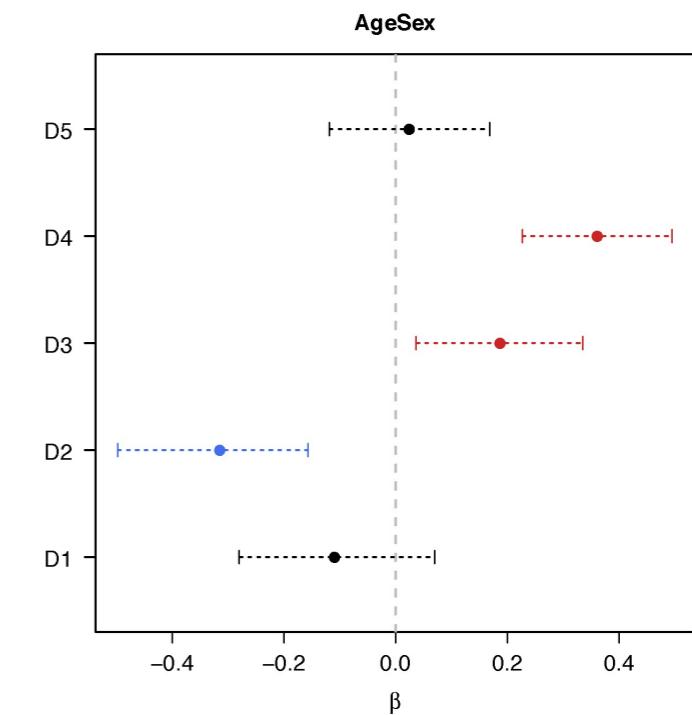
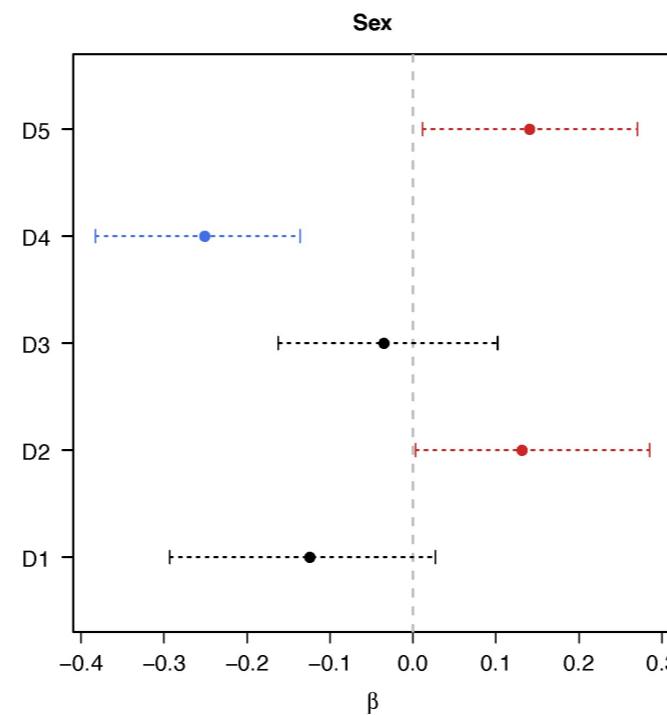
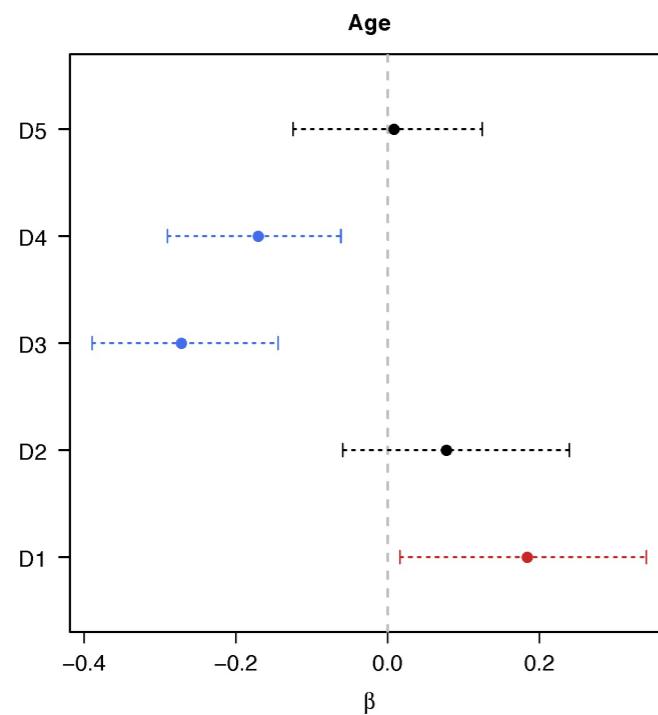
Table 1: Estimate (Est.) of β_1 , as well as standard error (SE), coverage probability with asymptotic variance in Theorem 1 (CP-A) and coverage probability from 500 bootstrap samples (CP-B) from different methods under the alternative hypothesis. All values are computed with $n = 100$ and $T_i = 100$ over 200 simulations.

Method	First Direction			Second Direction		
	Est. (SE)	CP-A	CP-B	Est. (SE)	CP-A	CP-B
Truth	-1.00	-	-	1.00	-	-
CAP	-1.00 (0.03)	0.950	0.950	0.81 (0.58)	0.885	0.870
CAP-OC	-1.00 (0.03)	0.950	0.950	0.52 (0.84)	0.730	0.715
CAP-C	-1.00 (0.03)	0.950	0.955	1.00 (0.03)	0.975	0.960
PCA	-0.02 (0.10)	-	0	-0.98 (0.03)	-	0
CPCA	-0.01 (0.11)	-	0	-1.00 (0.03)	-	0

PCA and common PCA do not find the first principal direction,
because they don't model covariates

Resting-state fMRI

Regression Coefficients



Age

Sex

Age*Sex

No statistical significant changes were found by massive edgewise regression

Brain Map of γ

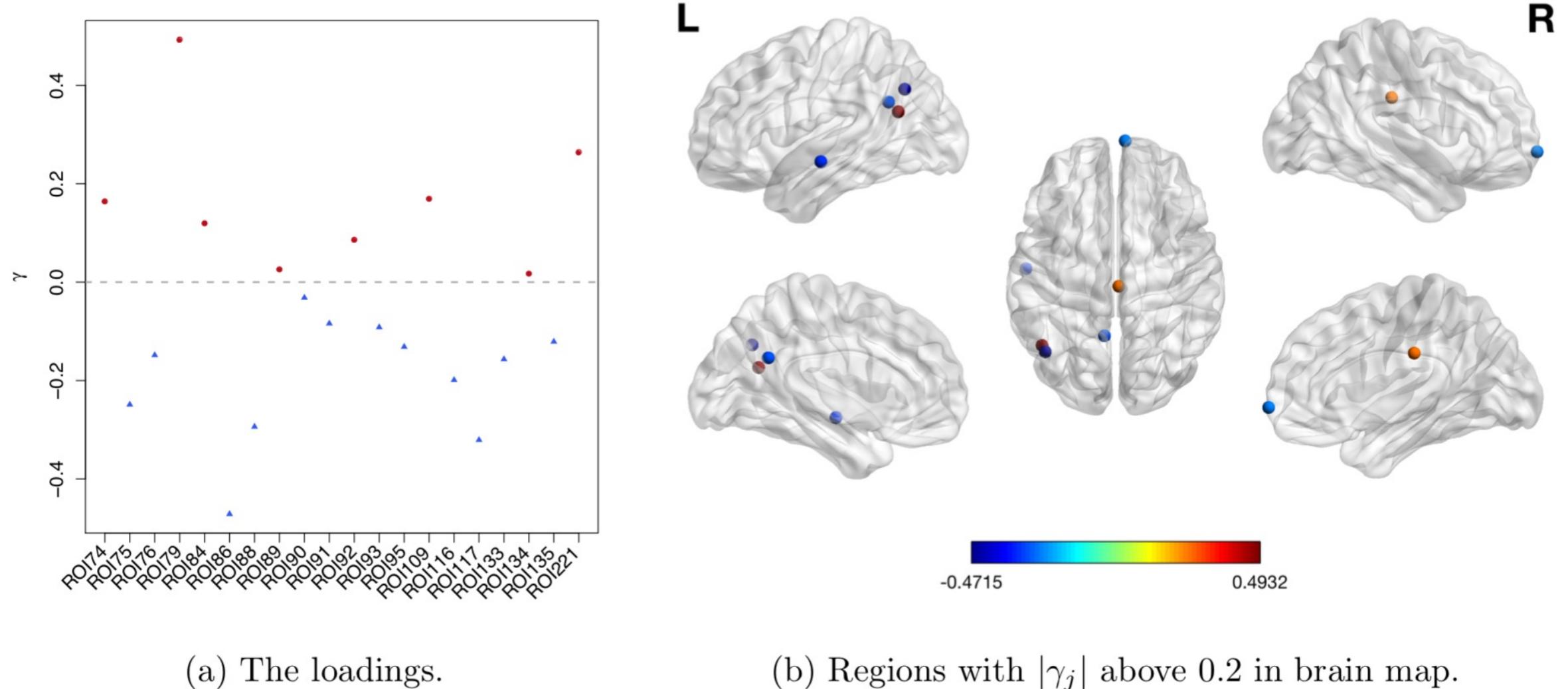


Figure 4: The loading profile and brain regions with absolute loading greater than 0.2 in projection direction D1 identified by CAP.

Discussion

- Regress **PD** matrices on vectors
- Method to identify covariate-related (supervised) directions vs (unsupervised) PCA
- Theorectical justifications
- Paper: Biostatistics ([10.1093/biostatistics/kxz057](https://doi.org/10.1093/biostatistics/kxz057))
- R pkg: **cap** 



Thank you!

Comments? Questions?

BigComplexData.com

or BrainDataScience.com

