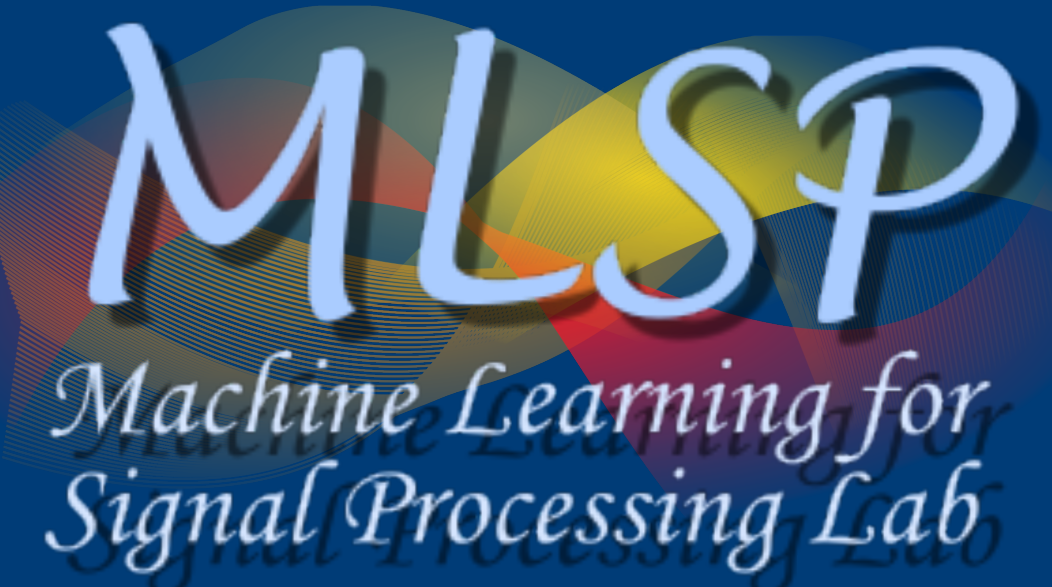


CONSTRAINED INDEPENDENT VECTOR ANALYSIS WITH REFERENCE FOR MULTI-SUBJECT FMRI ANALYSIS



Trung Vu¹ and Francisco Laport^{1,2} and Hanlu Yang¹ and Vince D. Calhoun³ and Tülay Adalı¹

¹ Department of Computer Science and Electrical Engineering, University of Maryland, Baltimore County, MD 21250, USA

² CITIC Research Center, University of A Coruña, Campus de Elviña, 15071 A Coruña, Spain

³ TReNDS, Georgia State University, Georgia Institute of Technology, and Emory University, Atlanta, GA 30303, USA



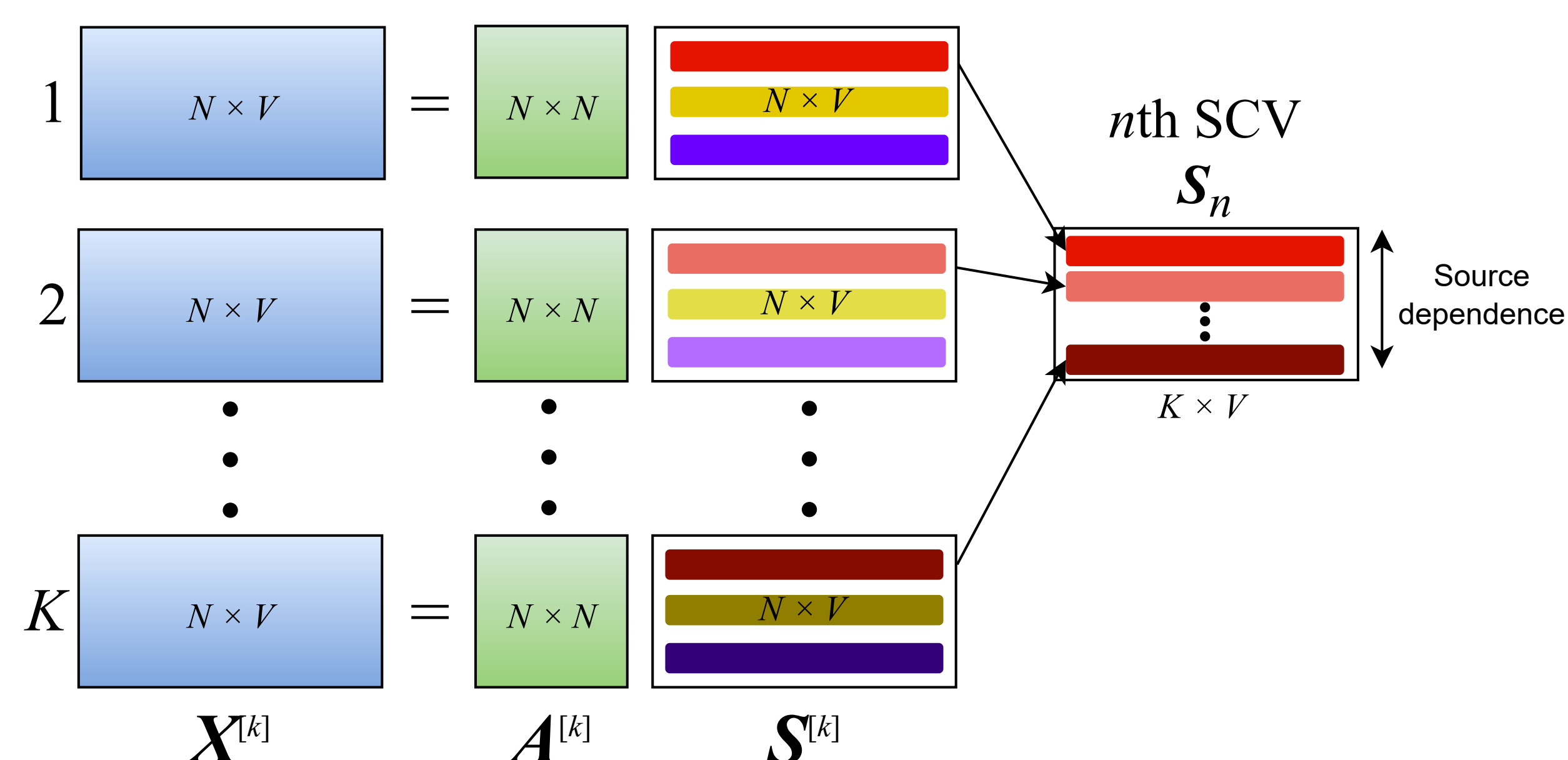
Introduction

- Independent vector analysis (IVA) is a joint blind source separation framework that exploits the **statistical dependencies across datasets**
- IVA has been successfully applied to various **neuroimaging** domains including multi-subject fMRI data analysis
- Constrained IVA (cIVA)** is an effective way to bypass *computational issues of IVA* and improve the quality of separation by incorporating available **prior information**
- Existing cIVA algorithms often rely on **user-defined threshold** values to define the constraints

Contributions

- Propose an adaptive-reverse scheme to select **variable thresholds** in cIVA, named ar-cIVA
- Propose a **threshold-free** formulation of cIVA by leveraging the unique structure of IVA, named tf-cIVA
- Show the **superior performance** of the two proposed algorithms compared with existing cIVA algorithms in different settings
- Demonstrate that they provide **meaningful and interpretable results** from analyzing real fMRI data

Independent Vector Analysis (IVA)

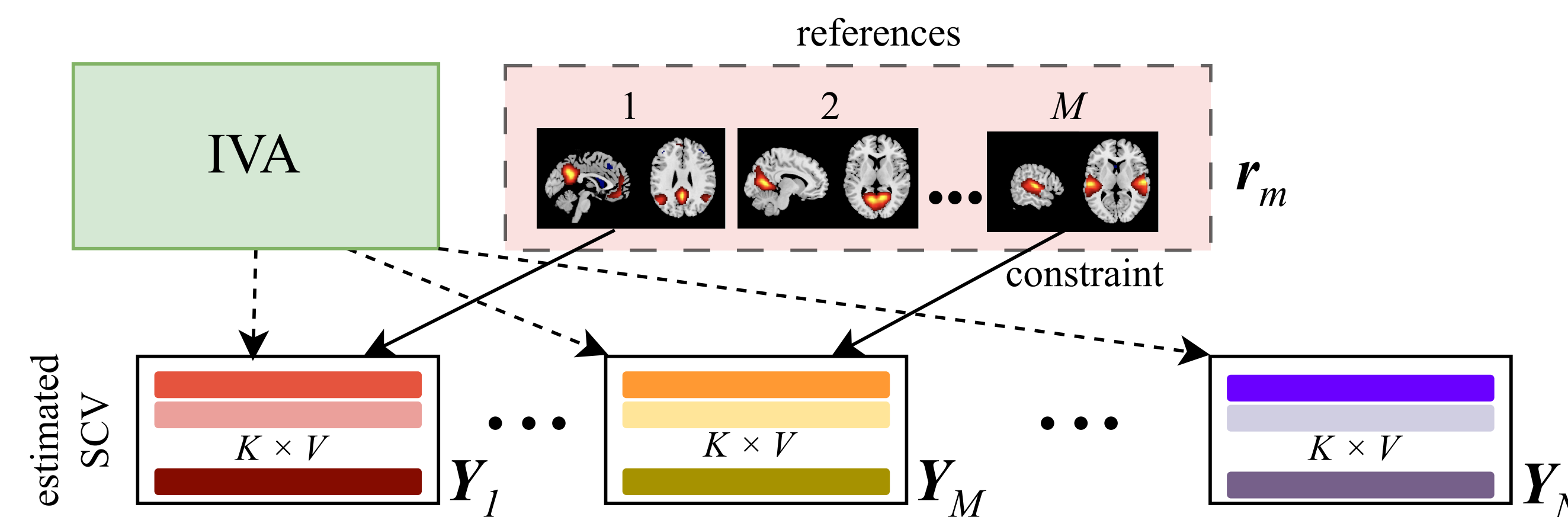


- The IVA cost function

$$\mathcal{J}_{\text{IVA}}(\mathbf{W}) \triangleq \sum_{n=1}^N \left(\sum_{k=1}^K \mathcal{H}(y_n^{[k]}) - \mathcal{I}(\mathbf{y}_n) \right) - \sum_{k=1}^K \log |\det(\mathbf{W}^{[k]})|$$

where $\mathcal{H}(y_n^{[k]})$ is the entropy of the n th estimated source for the k th dataset, $\mathcal{I}(\mathbf{y}_n)$ is the mutual information of the n th estimated source component vector (SCV), and $\mathbf{W}^{[k]}$ is the k th demixing matrix

Constrained IVA (cIVA)



- Constrained formulation of IVA with M references ($M \leq N$)
- $$\min_{\mathbf{W}} \mathcal{J}_{\text{IVA}}(\mathbf{W}) \text{ s.t. } \epsilon(\mathbf{r}_m, \mathbf{y}_m^{[k]}) \geq \rho_m^{[k]} \quad \forall m = 1, \dots, M \text{ and } k = 1, \dots, K$$

Proposed Algorithms for Constrained IVA

Adaptive-Reverse Constrained IVA (ar-cIVA)

$$\min_{\mathbf{W}} \mathcal{J}_{\text{IVA}}(\mathbf{W}) + \frac{1}{2\gamma} \sum_{m,k} \left(\left(\max \left(0, \mu_m^{[k]} + \gamma(\rho_m^{[k]} - \epsilon(\mathbf{r}_m, \mathbf{y}_m^{[k]})) \right) \right)^2 - (\mu_m^{[k]})^2 \right)$$

- Augmented Lagrangian** method is used as a stable approach to constrained optimization
- Adaptive-reverse scheme** alternates between two principles to select an appropriate threshold for each component
 - choosing the **smallest** value that does not satisfy the constraint

$$\rho_n^{[k]} = \operatorname{argmin} \{ \rho \in \mathcal{P} \mid \rho > \epsilon(\mathbf{r}_n, \mathbf{y}_n^{[k]}) \}$$
 - choosing the **largest** value that satisfies the constraint

$$\rho_n^{[k]} = \operatorname{argmax} \{ \rho \in \mathcal{P} \mid \rho \leq \epsilon(\mathbf{r}_n, \mathbf{y}_n^{[k]}) \}$$

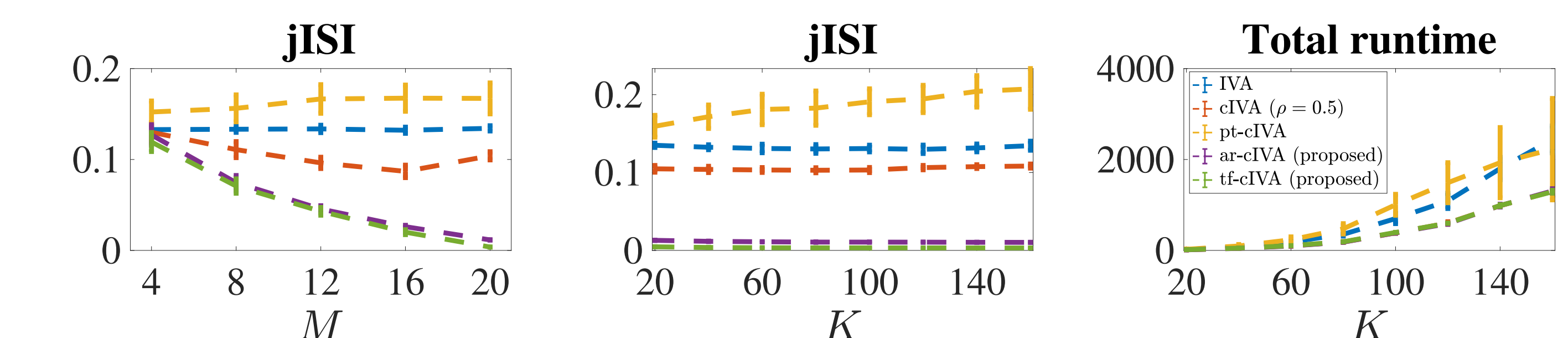
Threshold-Free Constrained IVA (tf-cIVA)

$$\min_{\mathbf{W}} \mathcal{J}_{\text{IVA}}(\mathbf{W}) + \frac{\lambda}{2} \sum_{m=1}^M \sum_{k=1}^K \left(\sum_{\substack{n=1 \\ n \neq m}}^M \epsilon^2(\mathbf{r}_m, \mathbf{y}_n^{[k]}) - \epsilon^2(\mathbf{r}_m, \mathbf{y}_m^{[k]}) \right)$$

- Multi-objective optimization** adds a *regularization* cost to
 - maximize** the **similarity** between the reference and the **corresponding** estimated source component
 - maximize** the **dissimilarity** between the reference and all the **other** estimated source components
- Selecting an appropriate value for λ **balances the trade-off** between the IVA cost and the constraint-regularization cost

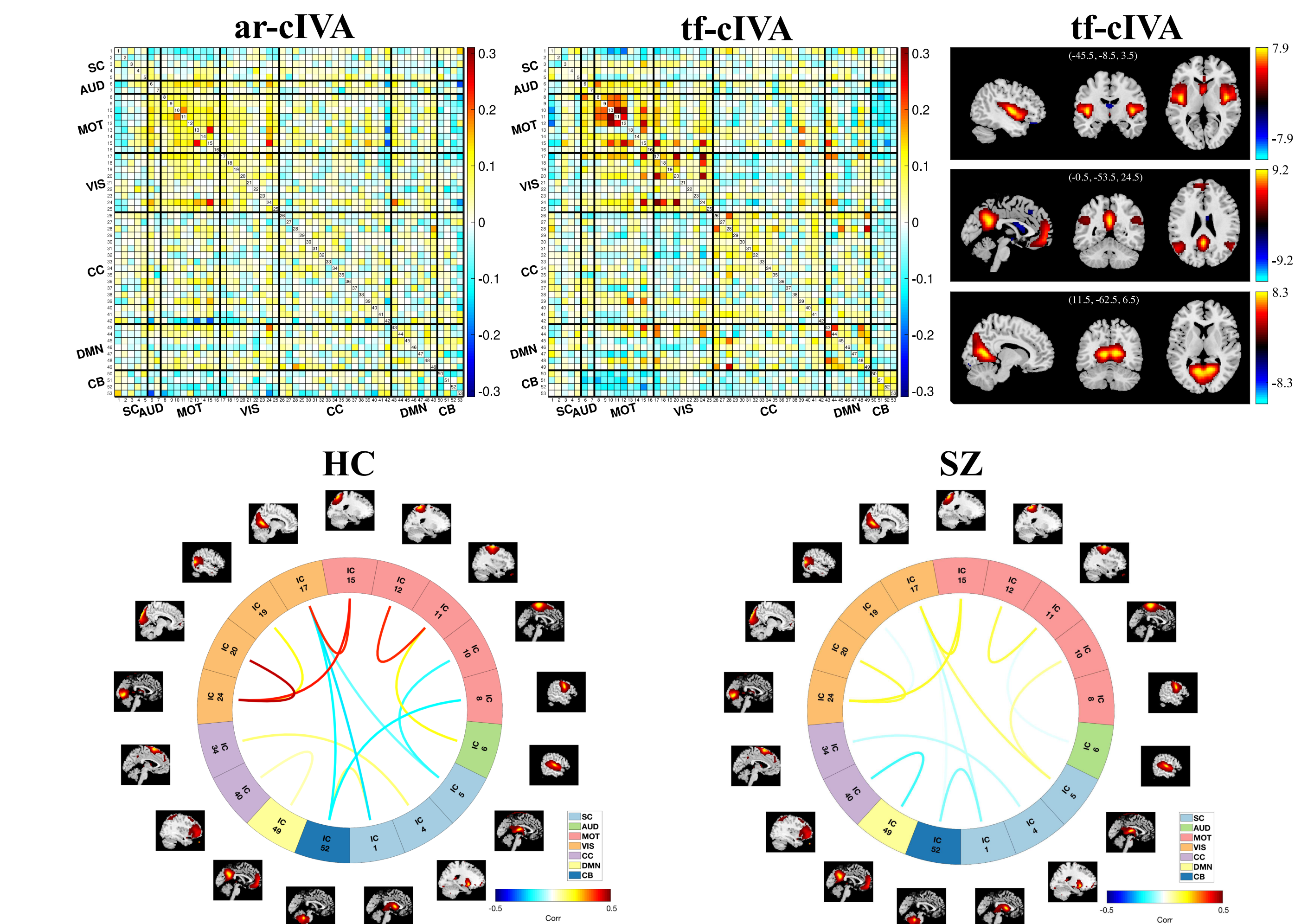
Experimental Results

Hybrid simulation — Varying M and K



- Two proposed cIVA algorithms **remarkably** outperform (unconstrained) IVA and existing cIVA algorithms
- tf-cIVA** slightly outperforms ar-cIVA

fMRI data analysis — $K = 98$ subjects



Summary

- tf-cIVA shows more **meaningful and interpretable results** when applied to real fMRI data
- tf-cIVA preserves **subject variability** and shows significant **group differences** between healthy control (HC) and schizophrenia patients (SZ)

This work is supported in part by the grants NIH R01MH118695, NIH R01MH123610, NIH R01AG073949, NSF 2316420, and Xunta de Galicia ED481B 2022/012.

