

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



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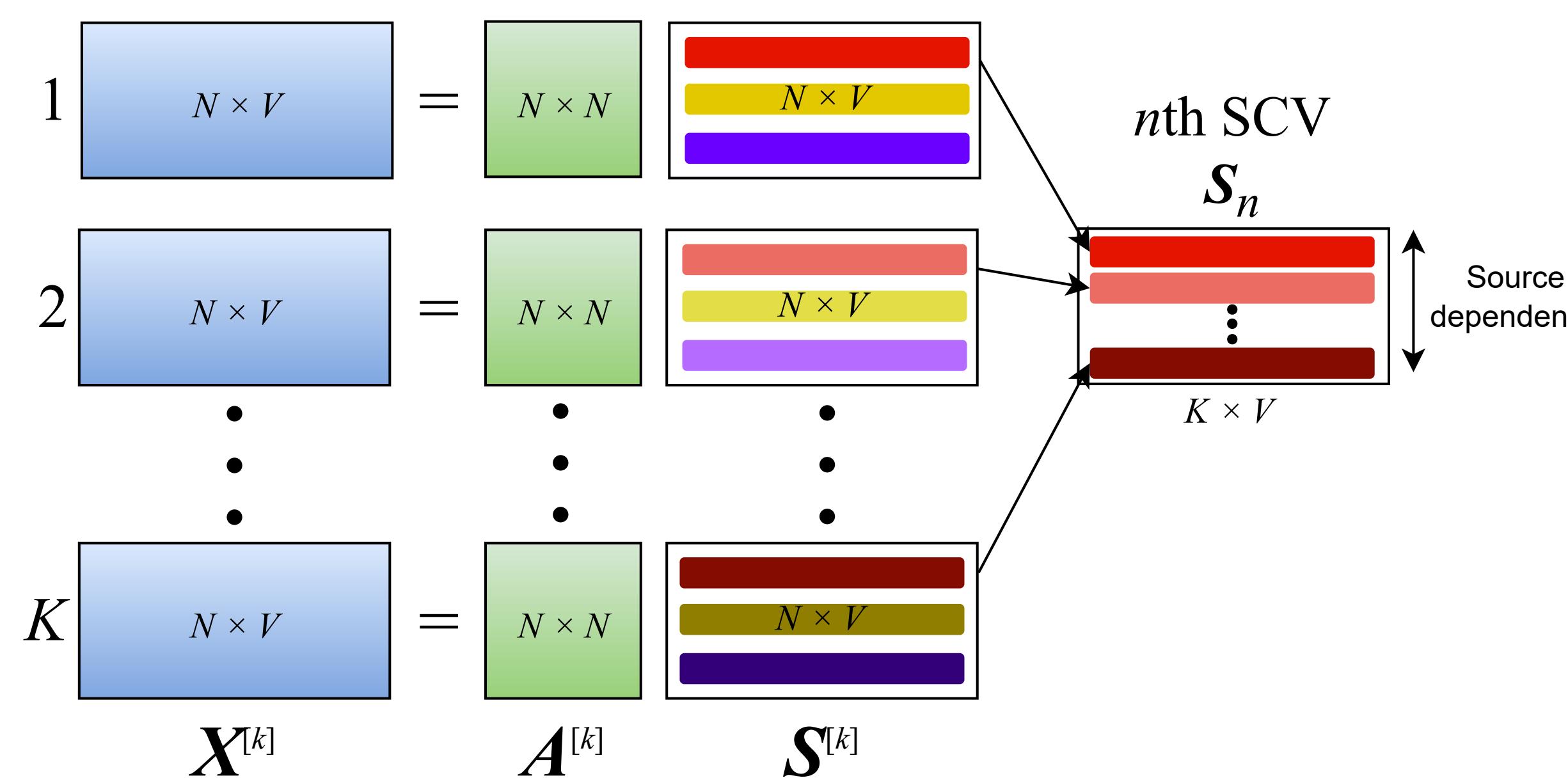
Introduction

- 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)

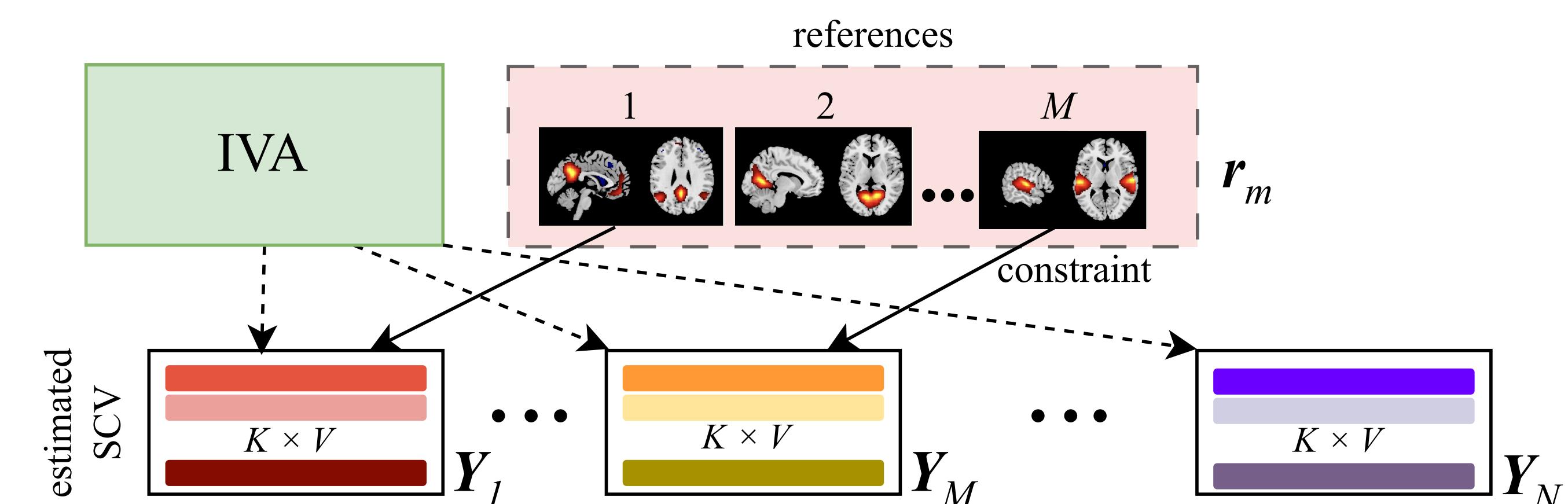


- The IVA cost function

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

where $\mathcal{H}(y_n^{[k]})$ is the entropy of the n th estimated source for the k th dataset, $\mathcal{I}(y_n)$ 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 ($M \leq N$)

$$\min_{\mathbf{W}} \mathcal{J}_{\text{IVA}}(\mathbf{W}) \text{ s.t. } \epsilon(r_m, y_m^{[k]}) \geq \rho_m^{[k]} \quad \forall m = 1, \dots, M \text{ and } k = 1, \dots, K$$

Proposed Algorithms for Constrained IVA

Adaptive-Reverse Constrained IVA (ar-cIVA)

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

- 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
 $\rho_n^{[k]} = \operatorname{argmin}\{\rho \in \mathcal{P} \mid \rho > \epsilon(r_n, y_n^{[k]})\}$
 - choosing the **largest** value that satisfies the constraint
 $\rho_n^{[k]} = \operatorname{argmax}\{\rho \in \mathcal{P} \mid \rho \leq \epsilon(r_n, y_n^{[k]})\}$

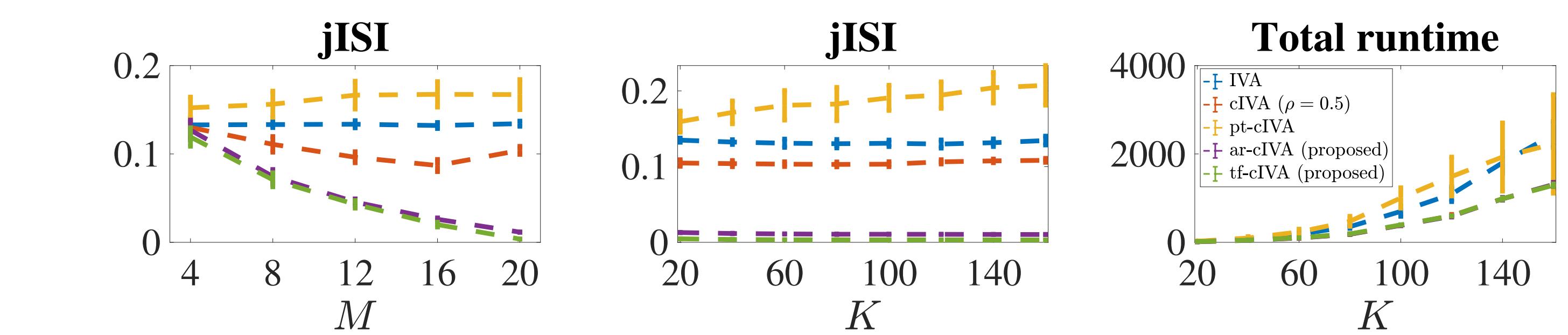
Threshold-Free Constrained IVA (tf-cIVA)

$$\min_{\mathbf{W}} \mathcal{J}_{\text{IVA}}(\mathbf{W}) + \frac{\lambda}{2} \sum_{m=1}^M \sum_{k=1}^K \left(\sum_{n=1}^M \epsilon^2(r_m, y_n^{[k]}) - \epsilon^2(r_m, y_m^{[k]}) \right)$$

- 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

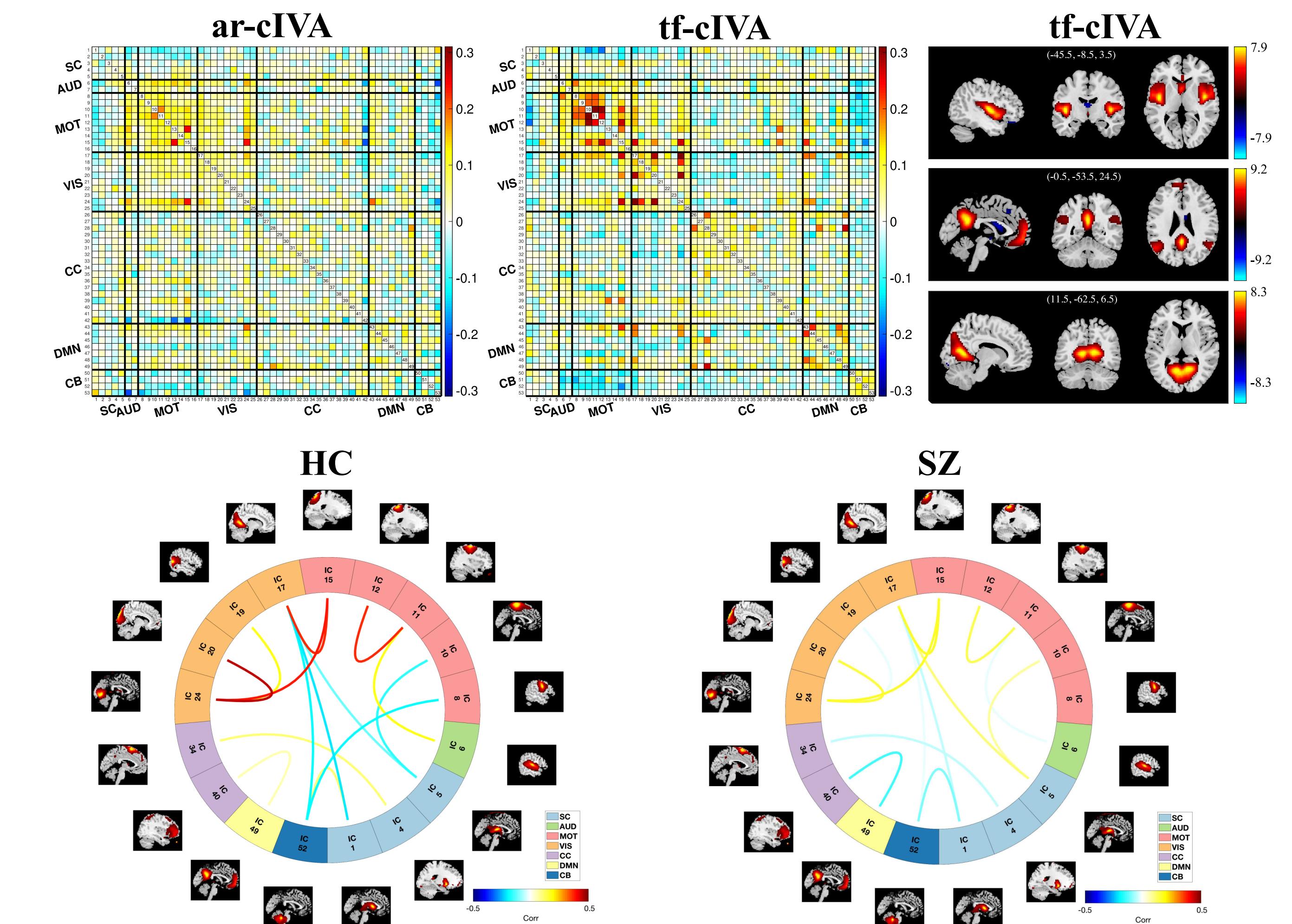
Experimental Results

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)

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