

# REPRODUCIBILITY IN JOINT BLIND SOURCE SEPARATION: APPLICATION TO FMRI ANALYSIS

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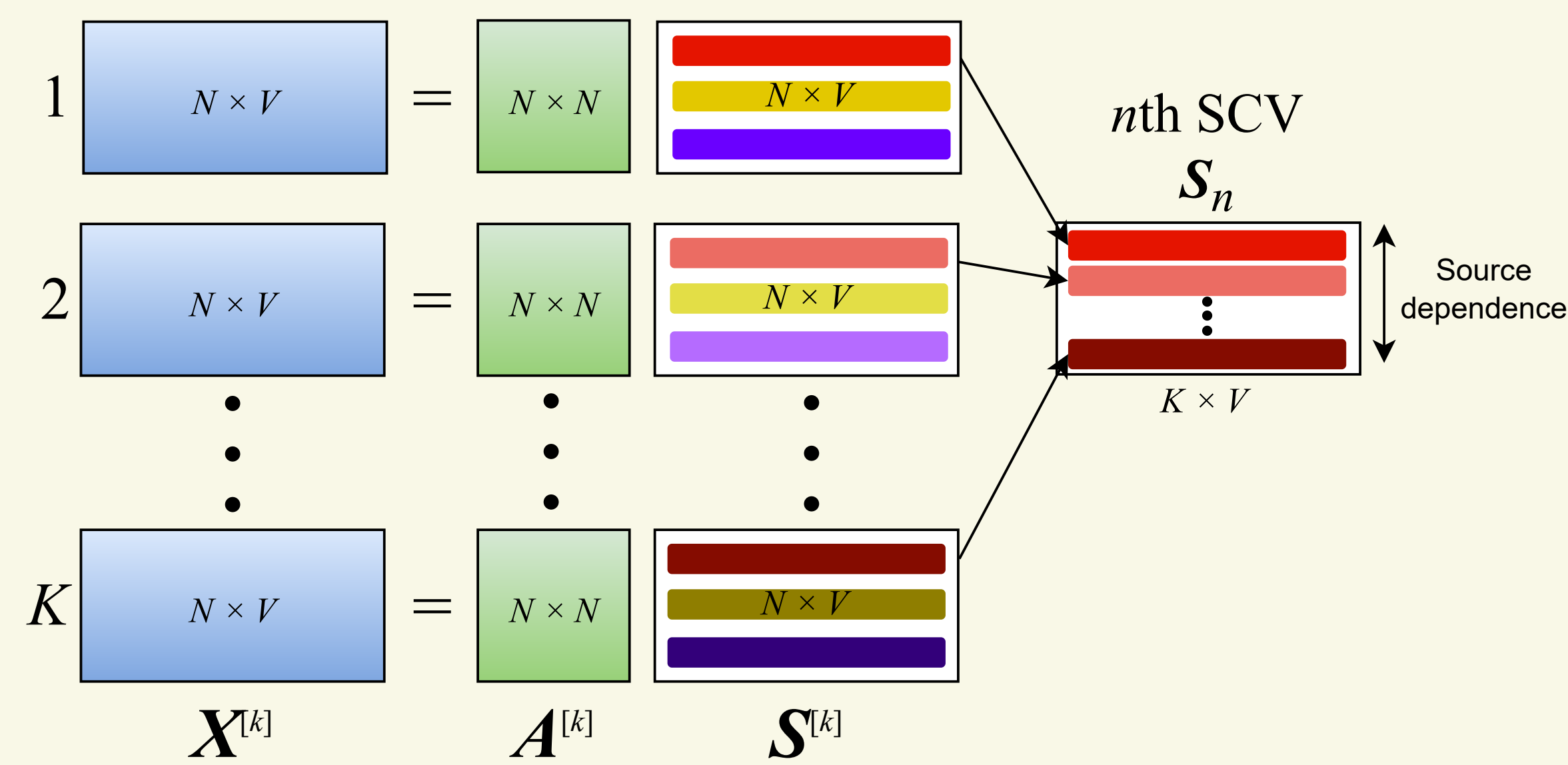
## Introduction

- ▶ Blind source separation (BSS) techniques have been successfully applied to a wide array of domains including fMRI data analysis
- ▶ Joint BSS (JBSS) techniques are implemented to leverage the joint information across multiple datasets
- ▶ JBSS results present high variability:
  - Cost functions of most JBSS algorithms are non-convex
  - Closed-form solutions do not exist for these problems – iterative solutions
  - No unique and perfect initialization – random initialization
- ▶ Reproducibility assessment of JBSS has been limited in the literature
- ▶ Highly consistent results do not guarantee a low bias in the estimates

## Contributions

- ▶ Evaluate the computational reproducibility of a JBSS algorithm: **constrained independent vector analysis (cIVA)**
- ▶ Propose a normalized measure related to the cost function to evaluate cIVA performance in **practical scenarios**
- ▶ Present a new mechanism for selecting the model complexity based on **reproducibility** and **accuracy** metrics
- ▶ Show that the model orders that balance accuracy and reproducibility metrics provide the most meaningful and interpretable results from analyzing real fMRI data

## Independent Vector Analysis (IVA)



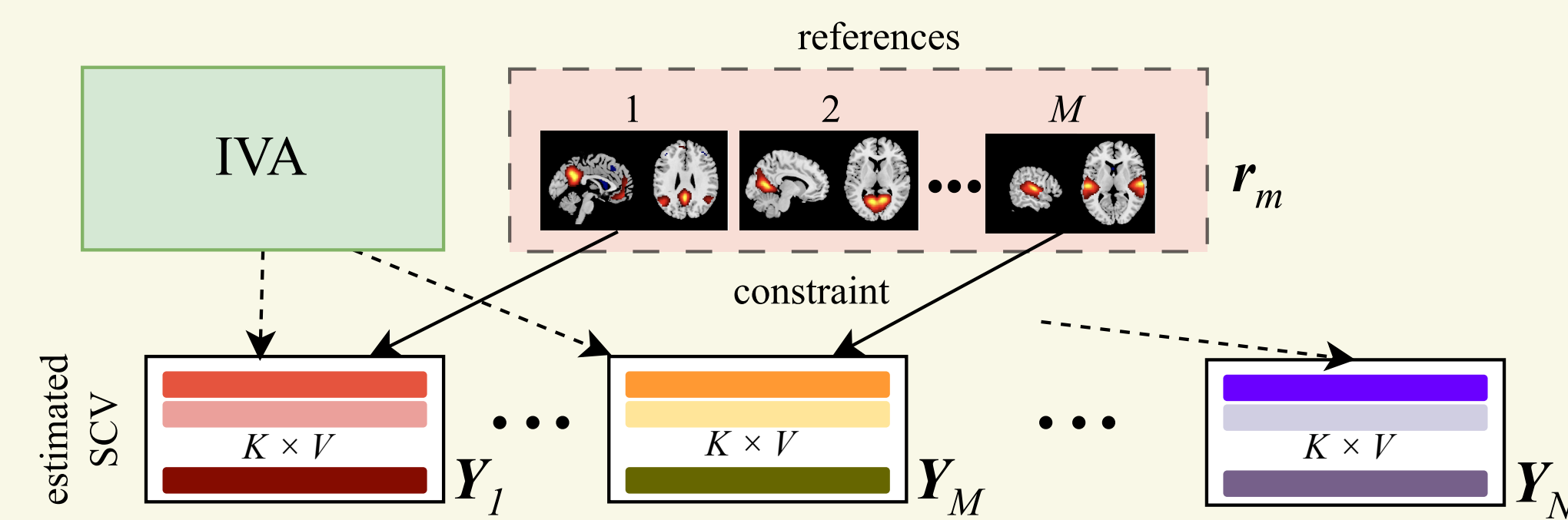
- ▶ IVA cost function

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

where  $\mathcal{W} = \{\mathbf{W}^{[1]}, \dots, \mathbf{W}^{[K]}\}$  are the demixing matrices of the  $K$  datasets,  $\mathbf{y}_n$  is the  $n$ th estimated SCV,  $\mathcal{I}(\mathbf{y}_n)$  is the mutual information of  $\mathbf{y}_n$  and  $\mathcal{H}(\mathbf{y}_n^{[k]})$  is the entropy of the  $n$ th estimated source for the  $k$ th dataset

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## Constrained IVA



- ▶ The augmented cost function

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

## Reproducibility and Accuracy Metrics

- ▶ In real applications the ground truth is unknown

- ▶ **Reproducibility: cross-joint-ISI**

- Let  $\mathbf{G}^{[k]} = \mathbf{A}^{[k]} \mathbf{W}^{[k]}$  and  $\mathbf{G} = 1/K \sum_{k=1}^K \mathbf{G}^{[k]}$  then

$$\text{ISI}(\mathbf{G}) = \frac{\sum_{i=1}^N \left( \frac{\sum_{j=1}^N |G_{ij}|}{\max_p |G_{ip}|} - 1 \right) + \sum_{j=1}^N \left( \frac{\sum_{i=1}^N |G_{ij}|}{\max_p |G_{jp}|} - 1 \right)}{2N(N-1)}$$

- Let  $\mathbf{W}_r^{[k]}$  be the  $k$ th demixing matrix of the  $r$ th run and  $\mathbf{P}_{ij} = 1/K \sum_{k=1}^K |(\mathbf{W}_i^{[k]})^{-1} \mathbf{W}_j^{[k]}|$  then

$$\text{cross-joint-ISI}_i = \frac{1}{R} \sum_{j=1, j \neq i}^R \text{ISI}(\mathbf{P}_{ij})$$

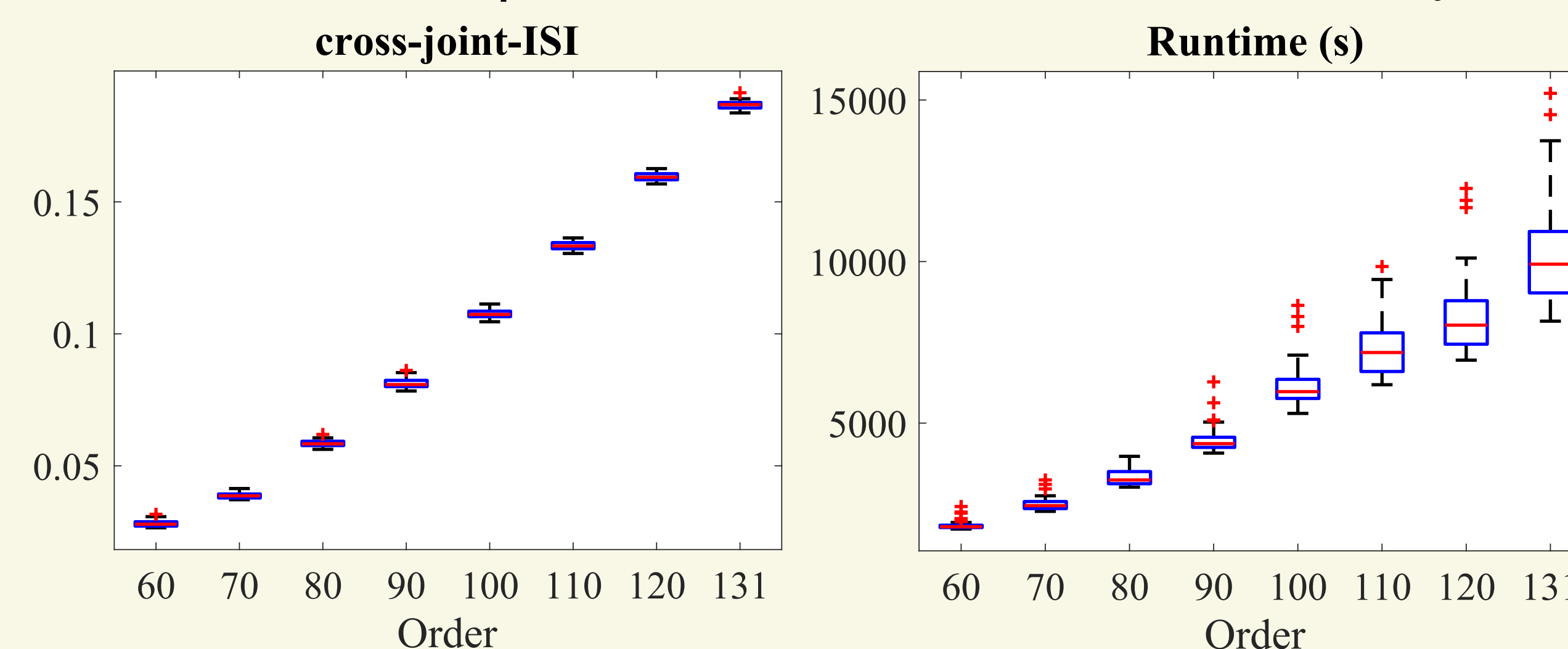
- ▶ **Accuracy: Pairwise normalized mutual information**

- Let  $\mathcal{I}(\mathbf{y}_i^{[k]}, \mathbf{y}_j^{[k]})$  be the mutual information between two estimated components

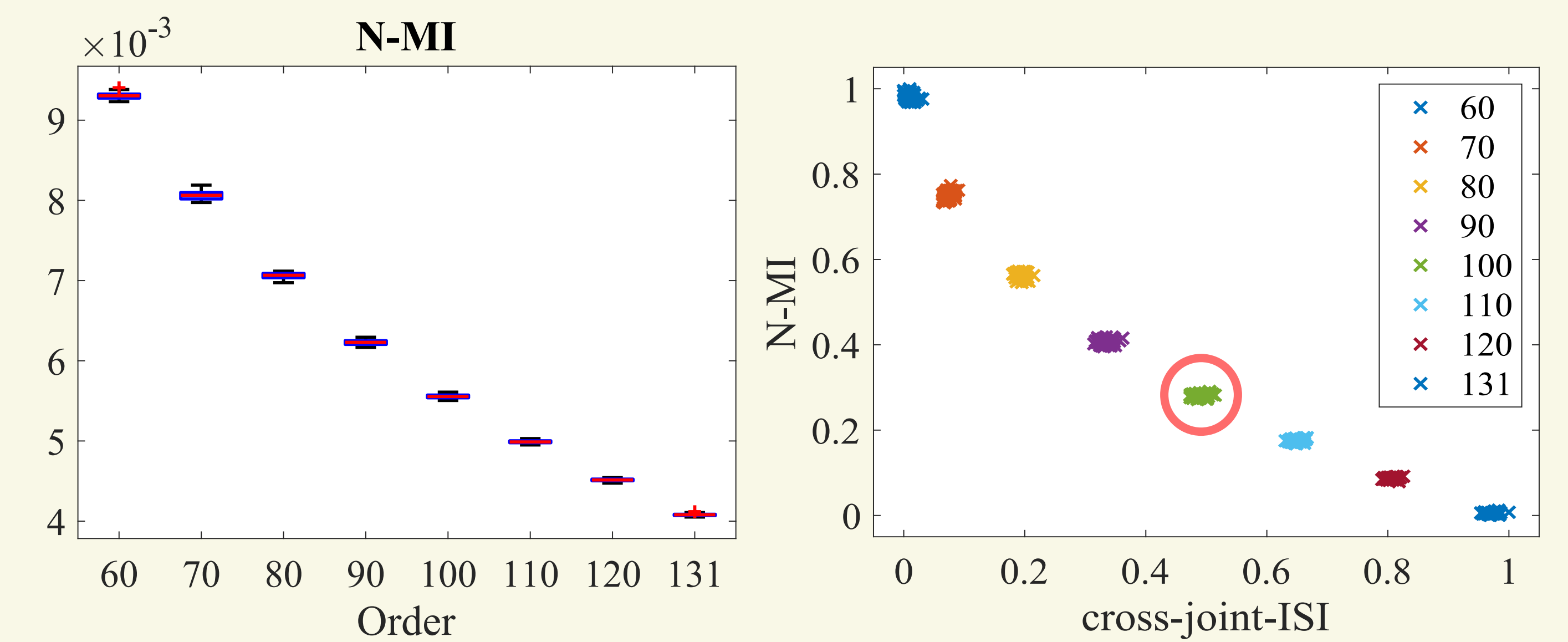
$$\text{N-MI}_r = \frac{2}{KN(N-1)} \sum_{k=1}^K \sum_{i=1, i \neq j}^N \sum_{j>i}^N \frac{2\mathcal{I}(\mathbf{y}_i^{[k]}, \mathbf{y}_j^{[k]})}{\mathcal{I}(\mathbf{y}_i^{[k]}, \mathbf{y}_i^{[k]}) + \mathcal{I}(\mathbf{y}_j^{[k]}, \mathbf{y}_j^{[k]})}$$

## Experimental Results

- ▶ Results from 50 independent runs on fMRI data from 98 subjects



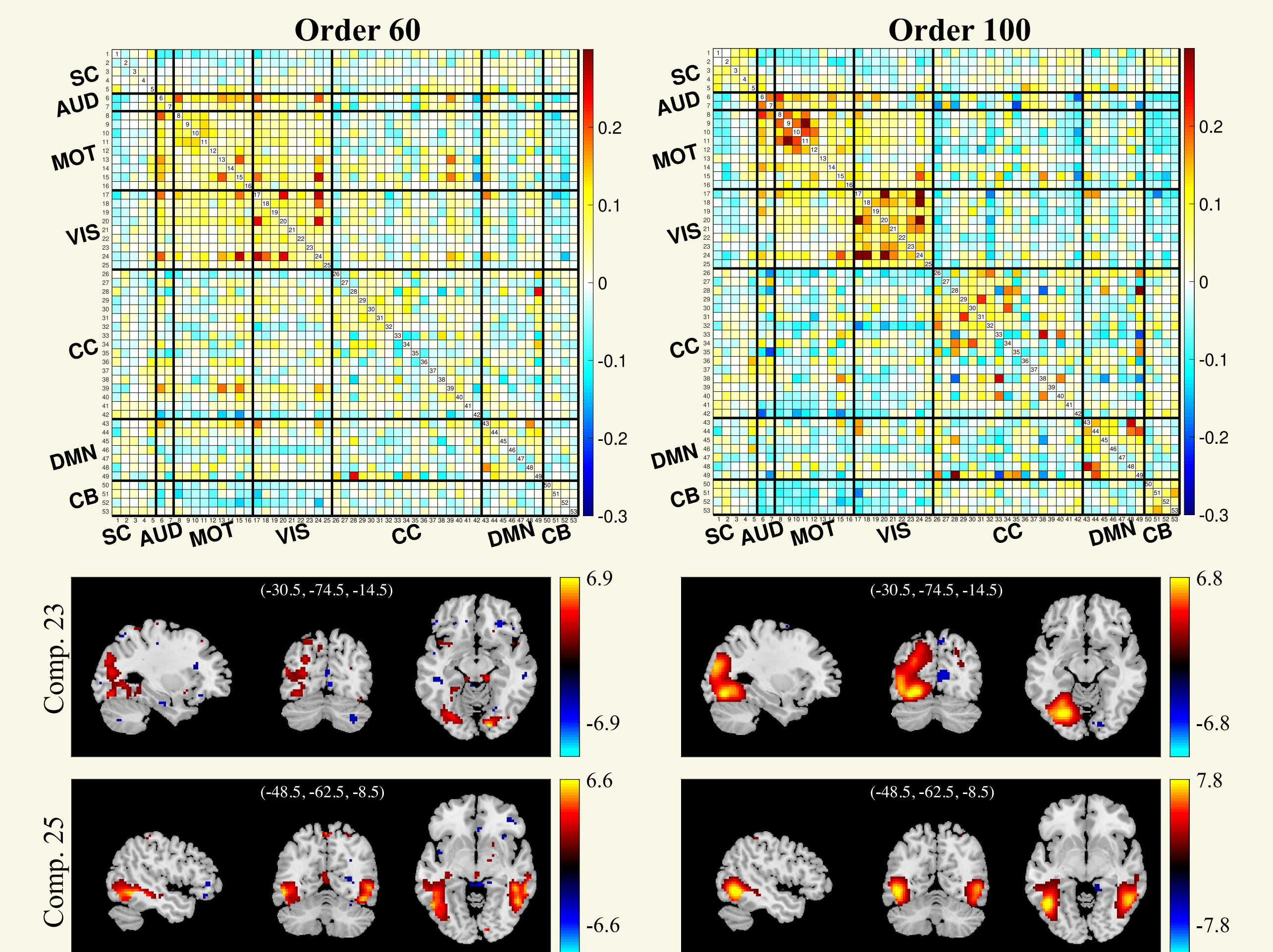
## Experimental Results (continued)



- ▶ Cross-joint-ISI increases with the model order while N-MI decreases – **Bias and Variance dilemma**
- ▶ Model orders balancing cross-joint-ISI and N-MI show **clearer spatial maps, higher temporal correlations** within functional domains, and **higher power ratios**

	60	80	90	100	120
<b>Power Ratio</b>	3.74±2	3.89±2.35	3.92±2.54	<b>4±2.82</b>	3.97±2.83

**Table:** Power ratio values for different model orders



## Summary

- ▶ Model order selection should be **guided** by both **reproducibility** and **accuracy** metrics
- ▶ Model orders that **balance the bias-variance tradeoff** provide a better model match and more interpretable and meaningful results in fMRI data