Shared Independent Component Analysis for Multi-Subject Neuroimaging

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https://github.com/hugorichard/ShICA

Problem

exposed to the same naturalistic stimuli (e.g. movie watching).

• Current well principled approaches need non-Gaussianity of

Solution: Shared ICA (ShICA)

ShICA performs multi-subjects ICA while modeling inter-subject

variability yielding an identifiable model even when common

sources are Gaussian. In practice, it yields better results than

ShICA

 $\mathbf{x}_i = A_i(\mathbf{s} + \mathbf{n}_i), i = 1, \dots, m$

Definition (Noise diversity): Let \mathcal{G} be the set of Gaussian

components. For all $j, j' \in \mathcal{G}, j \neq j'$, the sequences $(\Sigma_{ij})_{i=1...m}$ and

Theorem (Identifiability): Assuming noise diversity, ShICA is

Estimation by ShICA-ML

 $s_j \sim \frac{1}{2} \sum_{\alpha \in \{\frac{1}{2}, \frac{3}{2}\}} \mathcal{N}(0, \alpha)$

 $\mathbb{E}[s_j|\mathbf{x}] = \frac{\sum_{\alpha \in \{\frac{1}{2},\frac{3}{2}\}} \theta_{\alpha} h_{\alpha}}{\sum_{\alpha \in \{\frac{1}{2},\frac{3}{2}\}} \theta_{\alpha}}, \quad \mathbb{V}[s_j|\mathbf{x}] = \frac{\sum_{\alpha \in \{\frac{1}{2},\frac{3}{2}\}} \theta_{\alpha} g_{\alpha}}{\sum_{\alpha \in \{\frac{1}{2},\frac{3}{2}\}} \theta_{\alpha}}.$

M-step: Closed form updates for noise variances and quasi-Newton

Given m subjects, we model the data $\mathbf{x}_i \in \mathbb{R}^k$ of subject i as

• H1: s are independent components some of which may be

• H2: $\mathbf{n}_i \sim \mathcal{N}(0, \Sigma_i)$ where Σ_i is diagonal positive and \mathbf{n}_i

• H3: $\mathbb{E}[\mathbf{x}_i] = 0$, A_i invertible, $\mathbb{E}[\mathbf{s}\mathbf{s}^{\top}] = I_k$ and $m \geq 3$

 $(\Sigma_{ij'})_{i=1...m}$ are different where Σ_{ij} is the j, j entry of Σ_i .

identifiable up to a sign and permutation matrix.

Optimization Optimized via an EM algorithm.

We assume the ShICA model with:

updates for unmixing matrices.

• Uncover the shared neural responses of multiple subjects

the common sources to be identifiable.

competitive methods

Gaussian

E-step:

independent from s

Background (Multiset CCA): CCA of $(\mathbf{x}_i)_{i=1}^m$ given by solving $C\mathbf{u} = \lambda D\mathbf{u}$ where block i, j of C is $\mathbb{E}[\mathbf{x}_i \mathbf{x}_i^{\top}]$ and Dgiven by diagonal blocks of C.

Theorem (MultisetCCA solves ShICA: Assume x_i follows ShICA. Let $U = [\mathbf{u}_1 \dots \mathbf{u}_k]$ (first k eigenvectors of CCA problem) and $\lambda_1, \ldots, \lambda_k$ (first k eigenvalues). Set

Then if $\lambda_1 \dots \lambda_k$ are distinct, $W_i = P \Gamma_i A_i^{-1}$ where P is a permutation matrix and Γ_i a scaling matrix.

Serious problem:

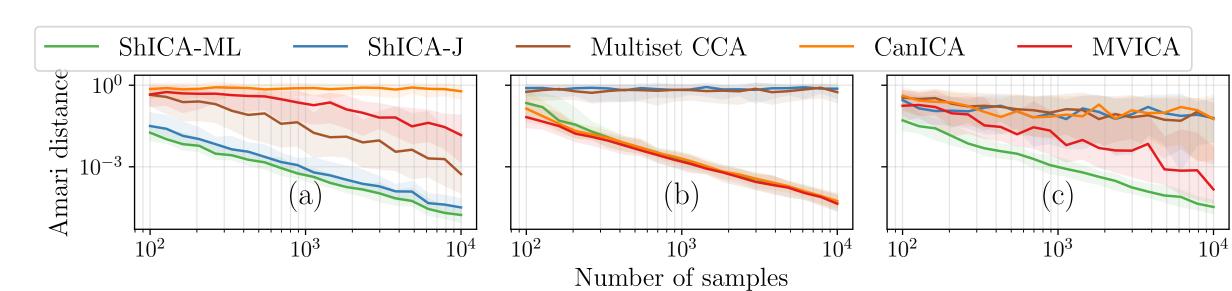
- The mapping from matrices to eigenvectors is highly non smooth
- ⇒ MultisetCCA fails in practice (especially when the eigenvalues are close).

span of the p leading eigenvectors is preserved: $W_i \approx QW_i$.

Last steps: Find scalings and noise variances (see paper)

Separation performance depending on the density of sources

Data generated with the ShICA model (m = 4 views, k = 5components). Non-Gaussian sources have Laplace densities.



Gaussian components with noise diversity (b) non-Gaussian components without noise diversity (c) Half of components are Gaussian with noise diversity, the other half is non-Gaussian without diversity

Estimation by ShICA-J

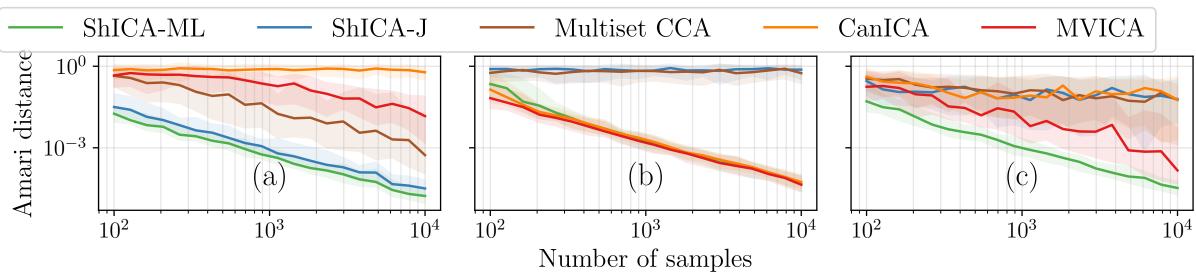
= U where $W_i \in \mathbb{R}^{k,k}$.

- We only have access to empirical covariance estimates

Solved by joint diagonalization:

- Large gap between the first k eigenvalues and other: the
- Recover Q by joint diagonalization of $Q\tilde{W}_{in}^{1}X_{i}X_{i}^{T}\tilde{W}_{i}^{T}Q^{T}$

Related work: SRM [2], MVICA [5], CanICA [6], IVA [1].

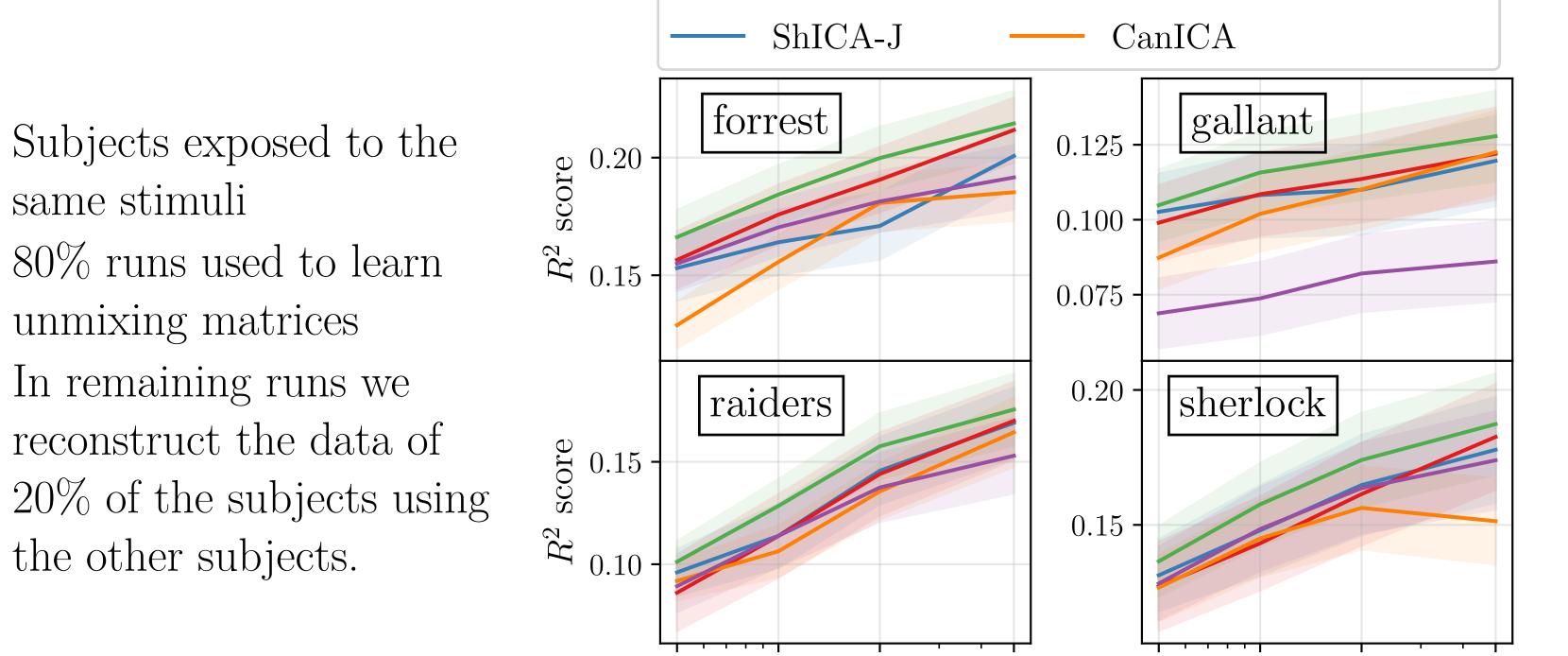






Reconstruction experiment fMRI

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MEG Phantom experiment

- 8 dipoles in a plastic head at different locations separately emit signal S_{true} during nepochs
- 20 sources are estimated: the best one is compared with S_{true} .

Subjects exposed to the

• 80% runs used to learn

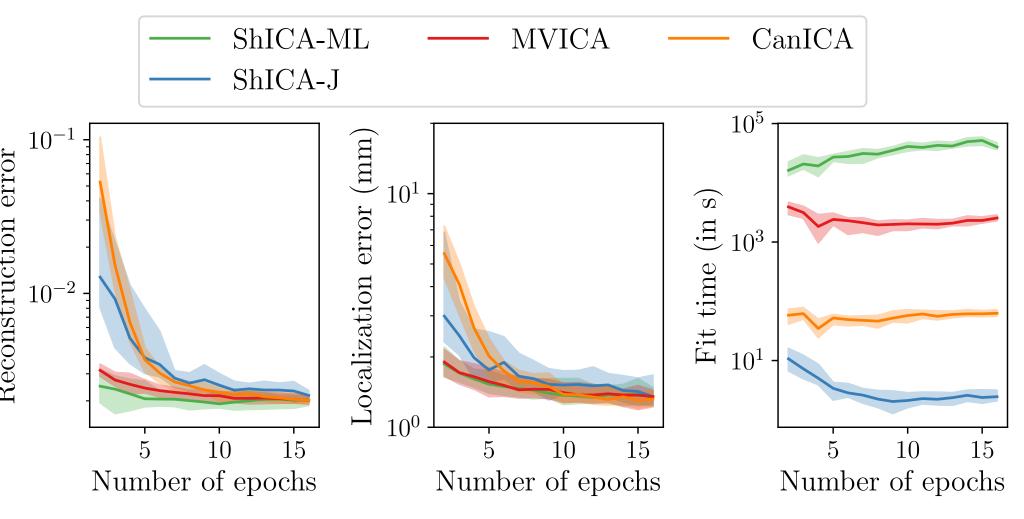
unmixing matrices

the other subjects.

In remaining runs we

reconstruct the data of

same stimuli



Number of components

References

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