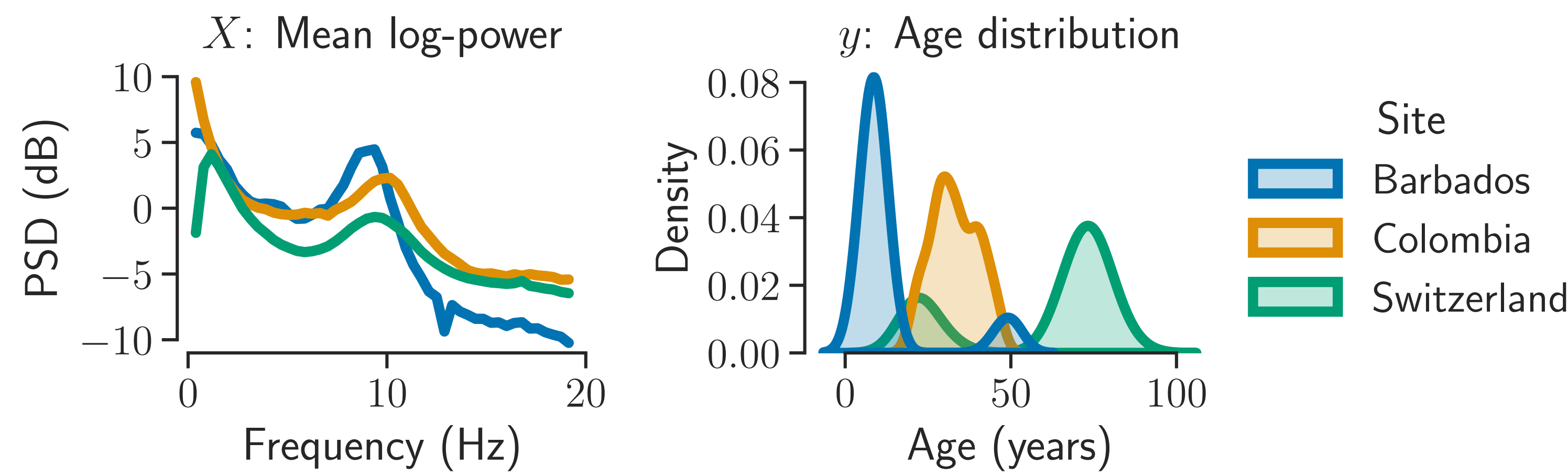




Multi-source domain adaptation

- **Distribution shift** between multiple source and target domains
- Domains in **neuroscience data**: subjects, recording sessions, experimental conditions, hospitals, acquisition protocols, ...



- In neuroscience: extensive use of **covariance matrices** that belong to the **Riemannian manifold**:

$$\mathbb{S}_d^{++} = \{\Sigma \in \mathbb{R}^{d \times d} \mid \Sigma^\top = \Sigma, \text{ and } \mathbf{x}^\top \Sigma \mathbf{x} > 0, \text{ for all } \mathbf{x} \neq 0\}$$

Question: How can we handle joint shifts in both Riemannian features Σ and the label y ?

Machine learning on \mathbb{S}_d^{++}

Classical geometric tools:

- **Riemannian mean** of $\{\Sigma_i\}_{i=1}^N$:

$$\bar{\Sigma} \triangleq \arg \min_{\Sigma \in \mathbb{S}_d^{++}} \sum_{i=1}^N \left\| \log \left(\Sigma^{-1/2} \Sigma_i \Sigma^{-1/2} \right) \right\|_F^2$$
- **Logarithmic mapping** projects Σ' to the tangent (Euclidean) space at Σ :

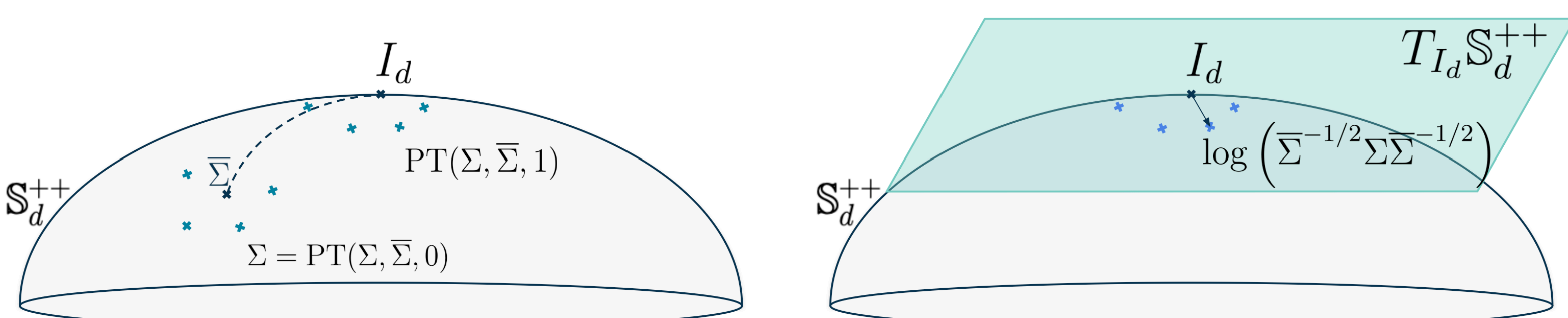
$$\log_{\Sigma}(\Sigma') \triangleq \Sigma^{1/2} \log \left(\Sigma^{-1/2} \Sigma' \Sigma^{-1/2} \right) \Sigma^{1/2} \in \mathbb{S}_d$$
- **Parallel transport** moves Σ' along the geodesic from Σ to I_d :

$$\text{PT}(\Sigma', \Sigma, \alpha) \triangleq \Sigma^{-\alpha/2} \Sigma' \Sigma^{-\alpha/2}$$

Learning from the dataset $\{(\Sigma_i, y_i)\}_{i=1}^N$ by transforming it into Euclidean vectors:

$$\phi(\Sigma_i, \bar{\Sigma}) \triangleq \text{uvec} \left(\log_{I_d} \left(\text{PT}(\Sigma_i, \bar{\Sigma}, 1) \right) \right) = \text{uvec} \left(\log \left(\bar{\Sigma}^{-1/2} \Sigma_i \bar{\Sigma}^{-1/2} \right) \right) \in \mathbb{R}^{d(d+1)/2}$$

where uvec vectorizes the upper triangular part [1]



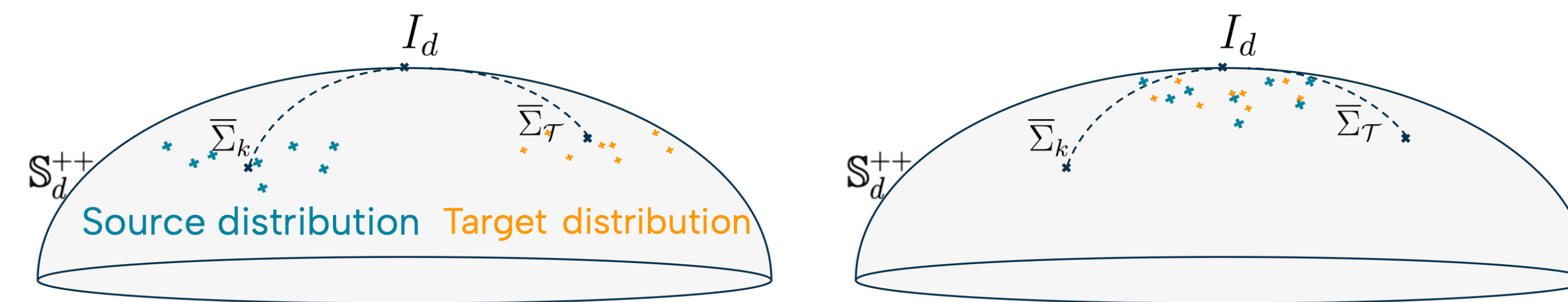
Multi-source domain adaptation on \mathbb{S}_d^{++}

K source domains: $\{(\Sigma_{k,i}, y_{k,i})\}_{i=1}^{N_k}$ for all $k \in \llbracket 1, K \rrbracket$ and $N_S = \sum_{k=1}^K N_k$
 One target domain: $\{(\Sigma_{\mathcal{T},i}, y_{\mathcal{T},i})\}_{i=1}^{N_{\mathcal{T}}}$

Assumption: $\bar{\Sigma}_{\mathcal{T}}$ is known.

- **Recenter** per domain $\mathcal{D} \in \{1, \dots, K, \mathcal{T}\}$ [2, 3]:

$$\mathbf{Z}_{\mathcal{D}} \triangleq [\phi(\Sigma_{\mathcal{D},1}, \bar{\Sigma}_{\mathcal{D}}), \dots, \phi(\Sigma_{\mathcal{D},N_{\mathcal{D}}}, \bar{\Sigma}_{\mathcal{D}})]^\top \in \mathbb{R}^{N_{\mathcal{D}} \times d(d+1)/2}$$



Problem: If $\mathbb{P}(y)$ varies across domains, then "Recenter" performs poorly.

Geodesic Optimization for Predictive Shift Adaptation: GOPSA

Assumption: $\bar{\Sigma}_{\mathcal{T}}$ and $\bar{y}_{\mathcal{T}}$ are known.

- **Learn the parallel transport** $\alpha \in [0, 1]$ **per domain**:

$$\phi(\Sigma, \bar{\Sigma}, \alpha) \triangleq \text{uvec} \left(\log_{I_d} \left(\text{PT}(\Sigma, \bar{\Sigma}, \alpha) \right) \right) = \text{uvec} \left(\log \left(\bar{\Sigma}^{-\alpha/2} \Sigma \bar{\Sigma}^{-\alpha/2} \right) \right)$$

- Learnable **source data matrix** w.r.t. $\gamma \in \mathbb{R}^K$:

$$\mathbf{Z}_S(\gamma) \triangleq [\phi(\Sigma_{1,1}, \bar{\Sigma}_1, \sigma(\gamma_1)), \dots, \phi(\Sigma_{K,N_K}, \bar{\Sigma}_K, \sigma(\gamma_K))]^\top$$

where $\sigma(\gamma) \triangleq (1 + \exp(-\gamma))^{-1}$ is the sigmoid function.

Train-time:

$$\gamma_S^* \triangleq \arg \min_{\gamma \in \mathbb{R}^K} \left\| \mathbf{y}_S - \mathbf{Z}_S(\gamma) \beta_S^*(\gamma) \right\|_2^2$$

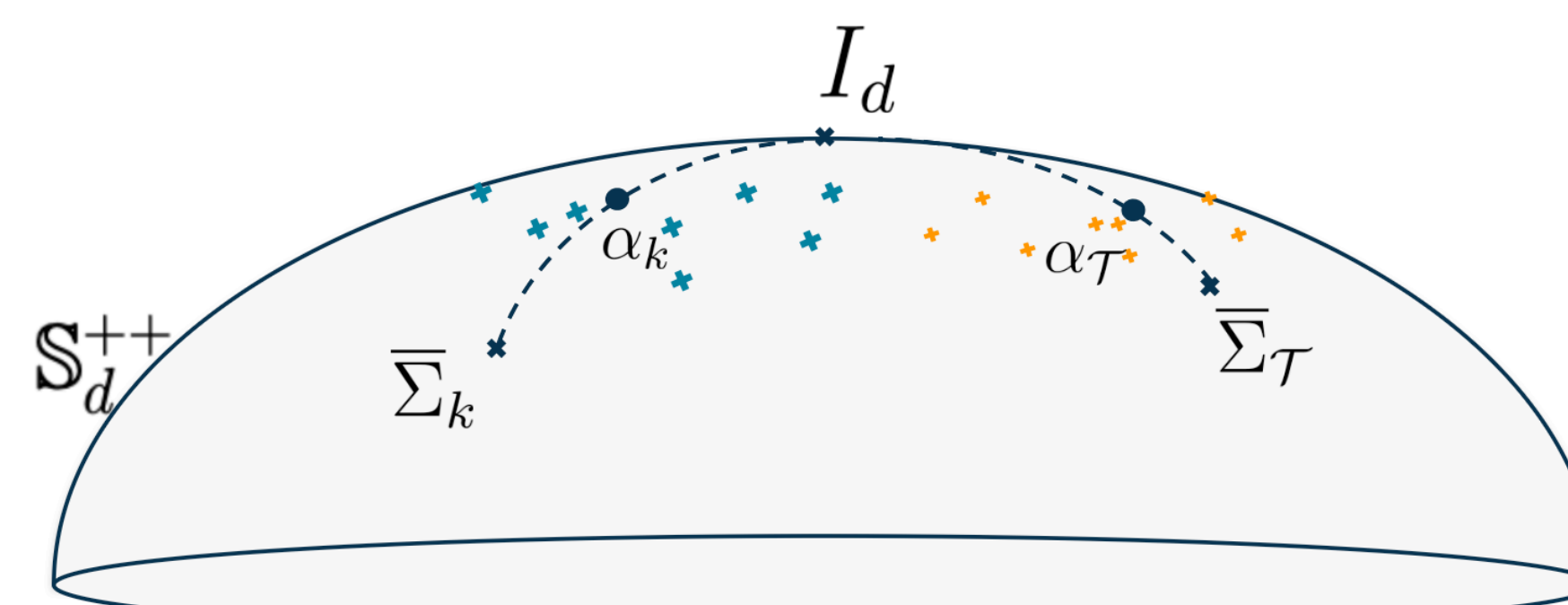
subject to $\beta_S^*(\gamma) \triangleq \mathbf{Z}_S(\gamma)^\top (\lambda \mathbf{I}_N + \mathbf{Z}_S(\gamma) \mathbf{Z}_S(\gamma)^\top)^{-1} \mathbf{y}_S$

- Learnable **target data matrix** w.r.t. $\gamma \in \mathbb{R}$:

$$\mathbf{Z}_{\mathcal{T}}(\gamma) \triangleq [\phi(\Sigma_{\mathcal{T},1}, \bar{\Sigma}_{\mathcal{T}}, \sigma(\gamma)), \dots, \phi(\Sigma_{\mathcal{T},N_{\mathcal{T}}}, \bar{\Sigma}_{\mathcal{T}}, \sigma(\gamma))]^\top$$

Test-time:

$$\begin{aligned} \text{(Optimization)} \quad \gamma_{\mathcal{T}}^* &\triangleq \arg \min_{\gamma \in \mathbb{R}} \left(\bar{y}_{\mathcal{T}} - \frac{1}{N_{\mathcal{T}}} \mathbf{1}_{N_{\mathcal{T}}}^\top \mathbf{Z}_{\mathcal{T}}(\gamma) \beta_S^*(\gamma_S^*) \right)^2 \\ \text{(Prediction)} \quad \hat{y}_{\mathcal{T}} &\triangleq \mathbf{Z}_{\mathcal{T}}(\gamma_{\mathcal{T}}^*) \beta_S^*(\gamma_S^*) \end{aligned}$$



Experimental results

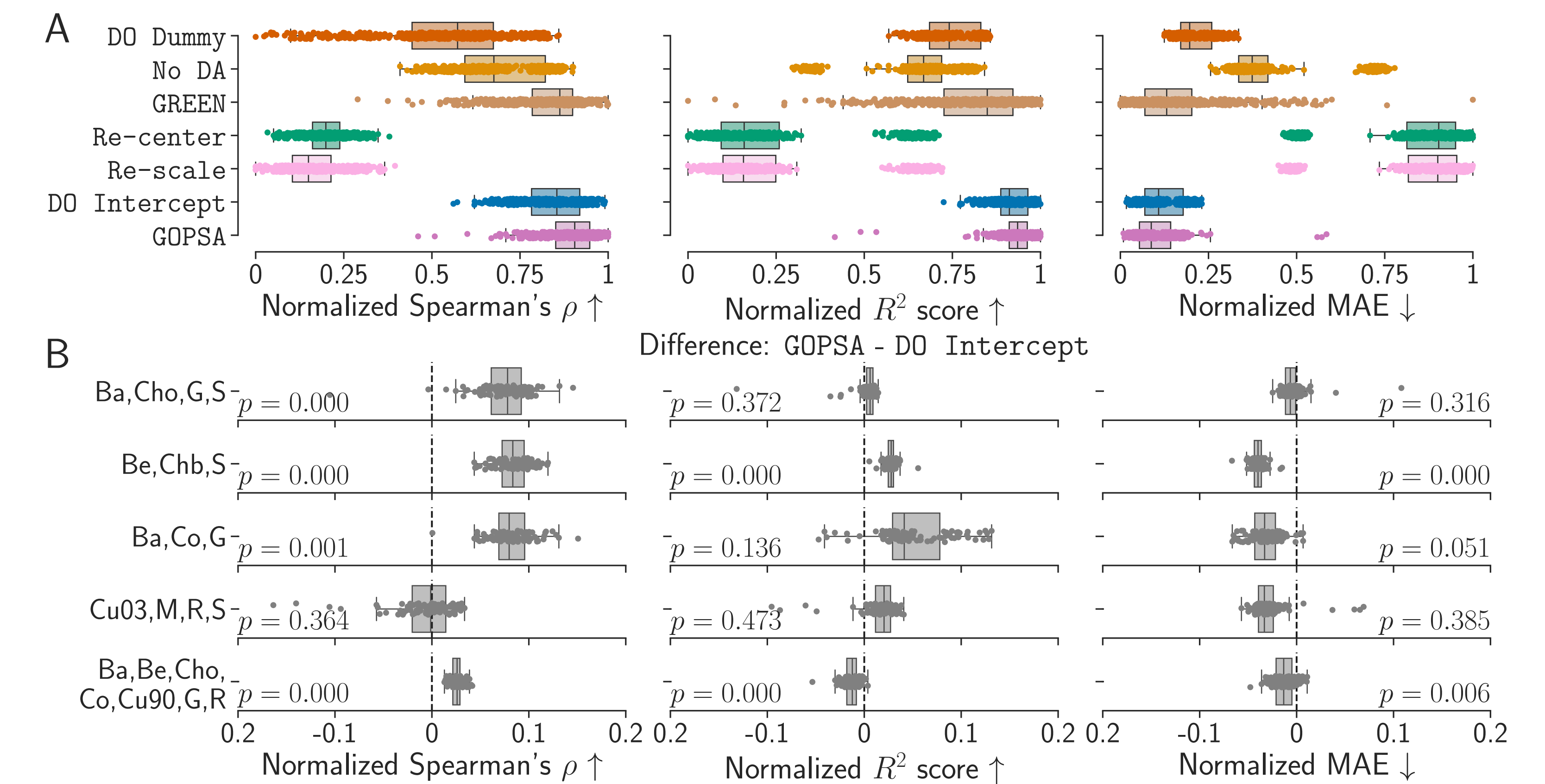
Benchmark on the HarMNqEEG dataset [4]:

- Brain age prediction task evaluated with 3 metrics: Spearman's ρ , R^2 , and MAE
- Cross power spectrum densities from 1500 participants with 49 frequency bins and 19 EEG sensors
- 5 source-site combinations from 11 sites: Barbados (Ba), Bern (Be), CHBMP Cuba (Chb), Columbia (Co), Chongqing (Cho), Cuba2003 (Cu03), Cuba90 (Cu90), Germany (G), Malaysia (M), Russia (R), and Switzerland (S)

Baselines:

- D0 Dummy: mean y per domain
- No DA: Ridge regression without domain adaptation
- GREEN [5]: SPDNet deep learning model
- Re-center/Re-scale [2, 3]: Ridge regression with a re-centering per domain
- D0 Intercept: Ridge regression with domain-specific bias

Results:



- Top performers: GOPSA and D0 Intercept, best average performance with low variance
- Paired t-test results: GOPSA outperformed D0 Intercept in 4 site combinations for Spearman's ρ , and 3 for MAE.

References

- [1] David Sabbagh et al. "Manifold-regression to predict from MEG/EEG brain signals without source modeling". In: *Advances in Neural Information Processing Systems* 32 (2019).
- [2] Paolo Zanini et al. "Transfer Learning: A Riemannian Geometry Framework With Applications to Brain-Computer Interfaces". In: *IEEE Transactions on Biomedical Engineering* 65.5 (May 2018), pp. 1107-1116. issn: 1558-2531. doi: 10.1109/TBME.2017.2742541.
- [3] Apolline Mellot et al. "Harmonizing and aligning M/EEG datasets with covariance-based techniques to enhance predictive regression modeling". In: *Imaging Neuroscience* 1 (2023), pp. 1-23.
- [4] Min Li et al. "Harmonized-Multinational qEEG norms (HarMNqEEG)". In: *NeuroImage* 256 (2022), p. 119190.
- [5] Joseph Paillard et al. "GREEN: a lightweight architecture using learnable wavelets and Riemannian geometry for biomarker exploration". In: *bioRxiv* (2024), pp. 2024-05.