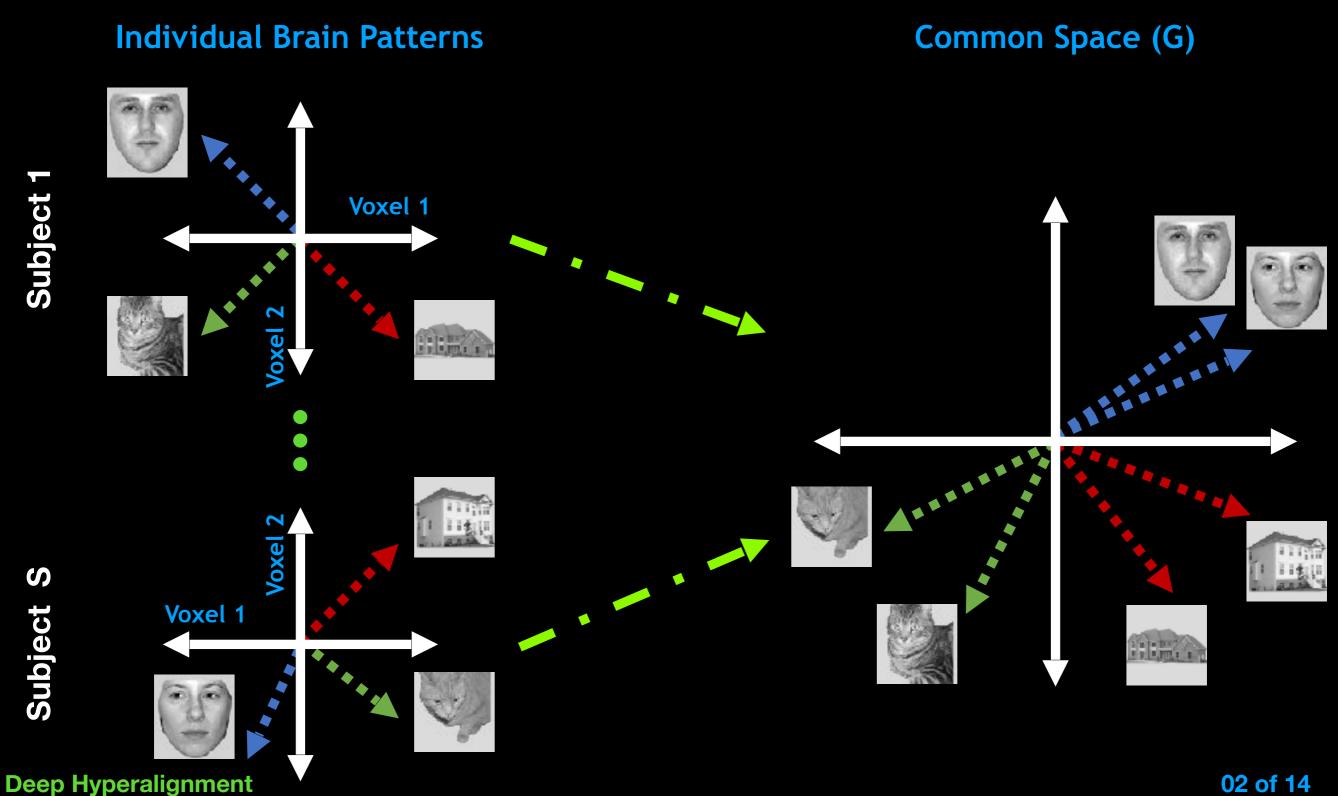


iBRAIN

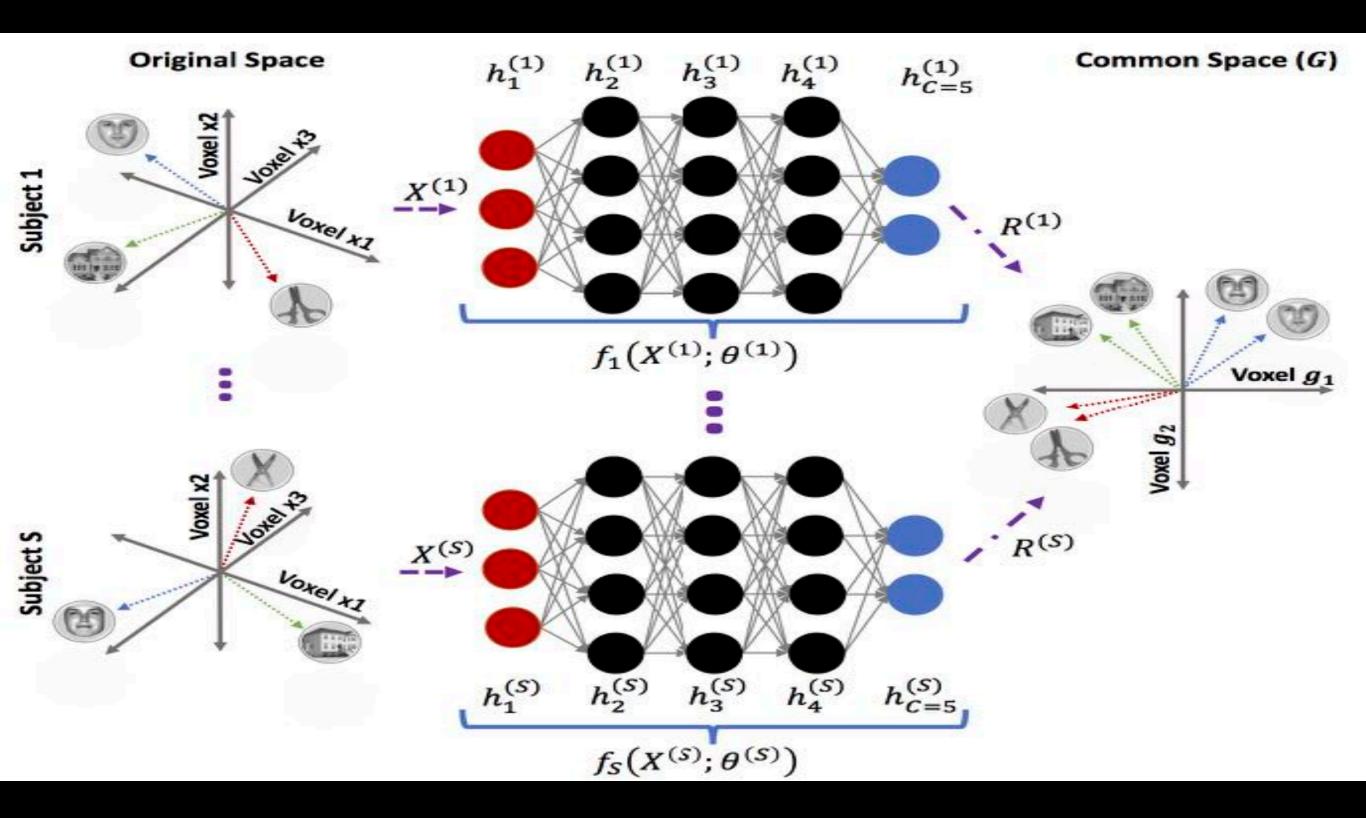
Deep Hyperalignment

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Hyperalignment



Main Idea



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DHA: Objective Function

★ We want to optimize following function:

$$\min_{\mathbf{G}, \mathbf{R}^{(i)}, \theta^{(i)}} \sum_{i=1}^{S} \left\| \mathbf{G} - f_i (\mathbf{X}^{(i)}; \theta^{(i)}) \mathbf{R}^{(i)} \right\|_F^2 \qquad \text{s.t.} \quad \mathbf{G}^\mathsf{T} \mathbf{G} = \mathbf{I}$$

$$\mathbf{G} = \frac{1}{S} \sum_{j=1}^{S} f_j(\mathbf{X}^{(j)}; \boldsymbol{\theta}^{(j)}) \mathbf{R}^{(j)}$$

where the deep network is defined as follows:

$$f_{\ell}(\mathbf{X}^{(\ell)}; \theta^{(\ell)}) = \mathsf{mat}\Big(\mathbf{h}_{C}^{(\ell)}, T, V_{new}\Big)$$

$$\mathbf{h}_m^{(\ell)} = \mathbf{g}\Big(\mathbf{W}_m^{(\ell)}\mathbf{h}_{m-1}^{(\ell)} + \mathbf{b}_m^{(\ell)}\Big), \quad \text{where} \quad \mathbf{h}_1^{(\ell)} = \mathbf{vec}\big(\mathbf{X}^{(\ell)}\big) \quad \text{and} \quad m = 2:C$$

DHA: Optimization

★ rank-m SVD

$$f_{\mathcal{C}}(\mathbf{X}^{(\ell)};\theta^{(\ell)}) \stackrel{SVD}{=} \mathbf{\Omega}^{(\ell)} \mathbf{\Sigma}^{(\ell)} (\mathbf{\Psi}^{(\ell)})^{\mathsf{T}}, \qquad \ell = 1:S$$

★ Projection Matrix

$$\mathbf{P}^{(\ell)} = f_{\ell} \left(\mathbf{X}^{(\ell)}; \boldsymbol{\theta}^{(\ell)} \right) \left(\left(f_{\ell} \left(\mathbf{X}^{(\ell)}; \boldsymbol{\theta}^{(\ell)} \right) \right)^{\mathsf{T}} f_{\ell} \left(\mathbf{X}^{(\ell)}; \boldsymbol{\theta}^{(\ell)} \right) + \epsilon \mathbf{I} \right)^{-1} \left(f_{\ell} \left(\mathbf{X}^{(\ell)}; \boldsymbol{\theta}^{(\ell)} \right) \right)^{\mathsf{T}}$$

$$= \mathbf{\Omega}^{(\ell)} \left(\mathbf{\Sigma}^{(\ell)} \right)^{\mathsf{T}} \left(\mathbf{\Sigma}^{(\ell)} \left(\mathbf{\Sigma}^{(\ell)} \right)^{\mathsf{T}} + \epsilon \mathbf{I} \right)^{-1} \mathbf{\Sigma}^{(\ell)} \left(\mathbf{\Omega}^{(\ell)} \right)^{\mathsf{T}} = \mathbf{\Omega}^{(\ell)} \mathbf{D}^{(\ell)} \left(\mathbf{\Omega}^{(\ell)} \mathbf{D}^{(\ell)} \right)^{\mathsf{T}}$$

where
$$\mathbf{D}^{(\ell)}(\mathbf{D}^{(\ell)})^{\mathsf{T}} = (\mathbf{\Sigma}^{(\ell)})^{\mathsf{T}}(\mathbf{\Sigma}^{(\ell)}(\mathbf{\Sigma}^{(\ell)})^{\mathsf{T}} + \epsilon \mathbf{I})^{-1}\mathbf{\Sigma}^{(\ell)}$$
.

Sum of Projection Matrices

$$\mathbf{A} = \sum_{i=1}^{S} \mathbf{P}^{(i)} = \widetilde{\mathbf{A}} \widetilde{\mathbf{A}}^{\mathsf{T}}, \quad \text{where} \quad \widetilde{\mathbf{A}} \in \mathbb{R}^{T \times mS} = \left[\mathbf{\Omega}^{(1)} \mathbf{D}^{(1)} ... \mathbf{\Omega}^{(S)} \mathbf{D}^{(S)} \right].$$
Decryptoperalignment

DHA: Optimization

★ Objective Function can be reformulated as follows:

$$\min_{\mathbf{G}, \mathbf{R}^{(i)}, \theta^{(i)}} \sum_{i=1}^{S} \left\| \mathbf{G} - f_i(\mathbf{X}^{(i)}; \theta^{(i)}) \mathbf{R}^{(i)} \right\| \equiv \max_{\mathbf{G}} \left(\operatorname{tr}(\mathbf{G}^{\mathsf{T}} \mathbf{A} \mathbf{G}) \right).$$

* So, we have:

AG = GA, where
$$\Lambda = \{\lambda_1 ... \lambda_T\}$$

$$\widetilde{\mathbf{A}} = \widetilde{\mathbf{G}} \widetilde{\boldsymbol{\Sigma}} \widetilde{\boldsymbol{\Psi}}^{\top} \longrightarrow \text{Incremental PCA}$$

★ DHA mappings can be calculated as follows:

$$\mathbf{R}^{(\ell)} = \left(\left(f_{\ell} \left(\mathbf{X}^{(\ell)}; \boldsymbol{\theta}^{(\ell)} \right) \right)^{\mathsf{T}} f_{\ell} \left(\mathbf{X}^{(\ell)}; \boldsymbol{\theta}^{(\ell)} \right) + \epsilon \mathbf{I} \right)^{-1} \left(f_{\ell} \left(\mathbf{X}^{(\ell)}; \boldsymbol{\theta}^{(\ell)} \right) \right)^{\mathsf{T}} \mathbf{G}.$$

DHA: Optimization

★ In order to use back-propagation algorithm for seeking an optimized parameters for the deep network, we also have:

$$\frac{\partial \mathbf{Z}}{\partial f_{\ell}(\mathbf{X}^{(\ell)};\theta^{(\ell)})} = 2\mathbf{R}^{(\ell)}\mathbf{G}^{\mathsf{T}} - 2\mathbf{R}^{(\ell)}(\mathbf{R}^{(\ell)})^{\mathsf{T}} \left(f_{\ell}(\mathbf{X}^{(\ell)};\theta^{(\ell)})\right)^{\mathsf{T}}.$$

where

$$\mathbf{Z} = \sum_{\ell=1}^{T} \lambda_{\ell}$$

Datasets

Table S2: The datasets.

Title	ID	S	K	T	V	X	Y	Z	Scanner	TR	TE
Mixed-gambles task	DS005	48	2	240	450	53	63	52	S 3T	2	30
Visual Object Recognition	DS105	71	8	121	1963	79	95	79	G 3T	2.5	30
Word and Object Processing	DS107	98	4	164	932	53	63	52	S 3T	2	28
Auditory and Visual Oddball	DS116	102	2	170	2532	53	63	40	P 3T	2	25
Multi-subject, multi-modal	DS117	171	2	210	524	64	61	33	S 3T	2	30
Forrest Gump	DS113	20	10	451	2400	160	160	36	S 7T	2.3	22
Raiders of the Lost Ark	N/A	10	7	924	980	78	78	54	S 3T	3	30

S is the number of subject; K denotes the number of stimulus categories; T is the number of scans in unites of TRs (Time of Repetition); V denotes the number of voxels in ROI; X, Y, Z are the size of 3D images; Scanners include S=Siemens, G = General Electric, and P = Philips in 3 Tesla or 7 Tesla; TR is Time of Repetition in millisecond; TE denotes Echo Time in second; Please see *openfmri.org* for more information.

Simple Tasks Analysis

Table 1: Accuracy of HA methods in post-alignment classification by using simple task datasets

\downarrow Algorithms, Datasets \rightarrow	DS005	DS105	DS107	DS116	DS117
ν-SVM [17]	71.65 ± 0.97	22.89 ± 1.02	38.84 ± 0.82	67.26 ± 1.99	73.32 ± 1.67
HA [1]	81.27 ± 0.59	30.03 ± 0.87	43.01 ± 0.56	74.23 ± 1.40	77.93 ± 0.29
RHA [2]	83.06 ± 0.36	32.62 ± 0.52	46.82 ± 0.37	78.71 ± 0.76	84.22 ± 0.44
KHA [3]	85.29 ± 0.49	37.14 ± 0.91	52.69 ± 0.69	78.03 ± 0.89	83.32 ± 0.41
SVD-HA [4]	90.82 ± 1.23	40.21 ± 0.83	59.54 ± 0.99	81.56 ± 0.54	95.62 ± 0.83
SRM [5]	91.26 ± 0.34	48.77 ± 0.94	64.11 ± 0.37	83.31 ± 0.73	95.01 ± 0.64
SL [9]	90.21 ± 0.61	49.86 ± 0.4	64.07 ± 0.98	82.32 ± 0.28	94.96 ± 0.24
CAE [6]	94.25 ± 0.76	54.52 ± 0.80	72.16 ± 0.43	91.49 ± 0.67	95.92 ± 0.67
DHA	97.92 ± 0.82	60.39 ± 0.68	73.05 ± 0.63	90.28 ± 0.71	97.99 ± 0.94

Table 2: Area under the ROC curve (AUC) of different HA methods in post-alignment classification by using simple task datasets

↓Algorithms, Datasets→	DS005	DS105	DS107	DS116	DS117
ν-SVM [17]	68.37±1.01	21.76 ± 0.91	36.84 ± 1.45	62.49±1.34	70.17 ± 0.59
HA [1]	70.32 ± 0.92	28.91 ± 1.03	40.21 ± 0.33	70.67 ± 0.97	76.14 ± 0.49
RHA [2]	82.22 ± 0.42	30.35 ± 0.39	43.63 ± 0.61	76.34 ± 0.45	81.54 ± 0.92
KHA [3]	80.91 ± 0.21	36.23 ± 0.57	50.41 ± 0.92	75.28 ± 0.94	80.92 ± 0.28
SVD-HA [4]	88.54 ± 0.71	37.61 ± 0.62	57.54 ± 0.31	78.66 ± 0.82	92.14 ± 0.42
SRM [5]	90.23 ± 0.74	44.48 ± 0.75	62.41 ± 0.72	79.20 ± 0.98	93.65 ± 0.93
SL [9]	89.79 ± 0.25	47.32 ± 0.92	61.84 ± 0.32	80.63 ± 0.81	93.26 ± 0.72
CAE [6]	91.24 ± 0.61	52.16 ± 0.63	72.33 ± 0.79	87.53 ± 0.72	91.49 ± 0.33
DHA	96.91±0.82	59.57±0.32	70.23 ± 0.92	89.93±0.24	96.13±0.32

Complex Tasks Analysis

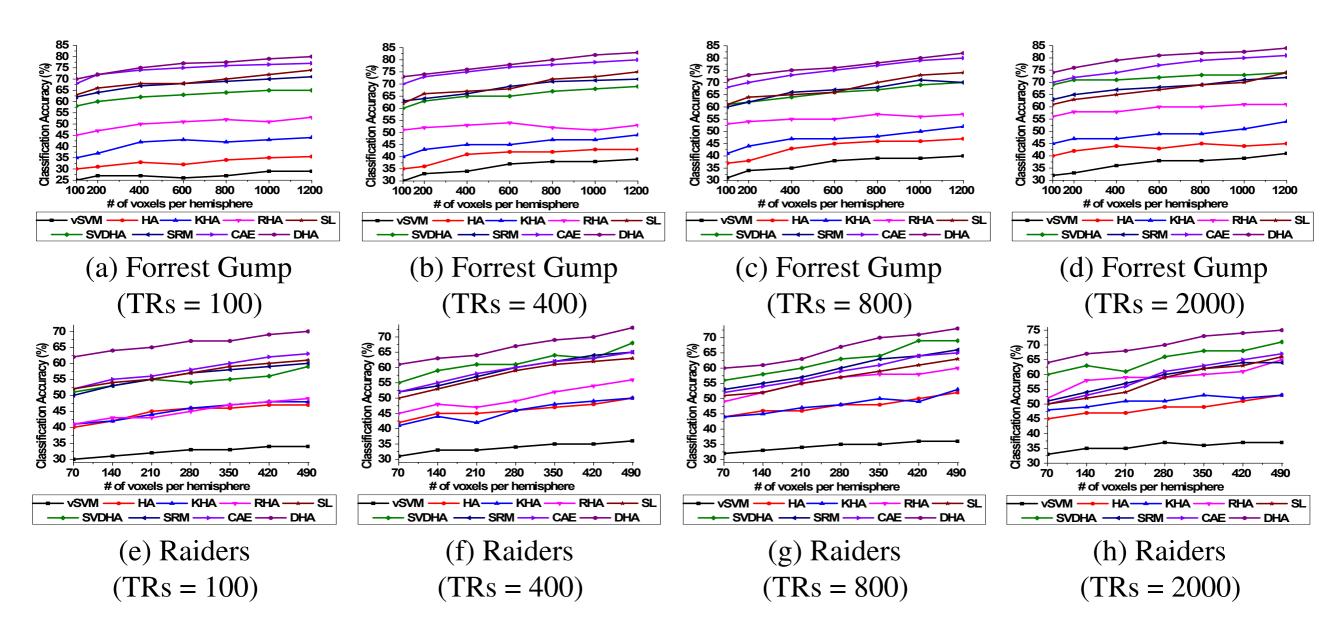
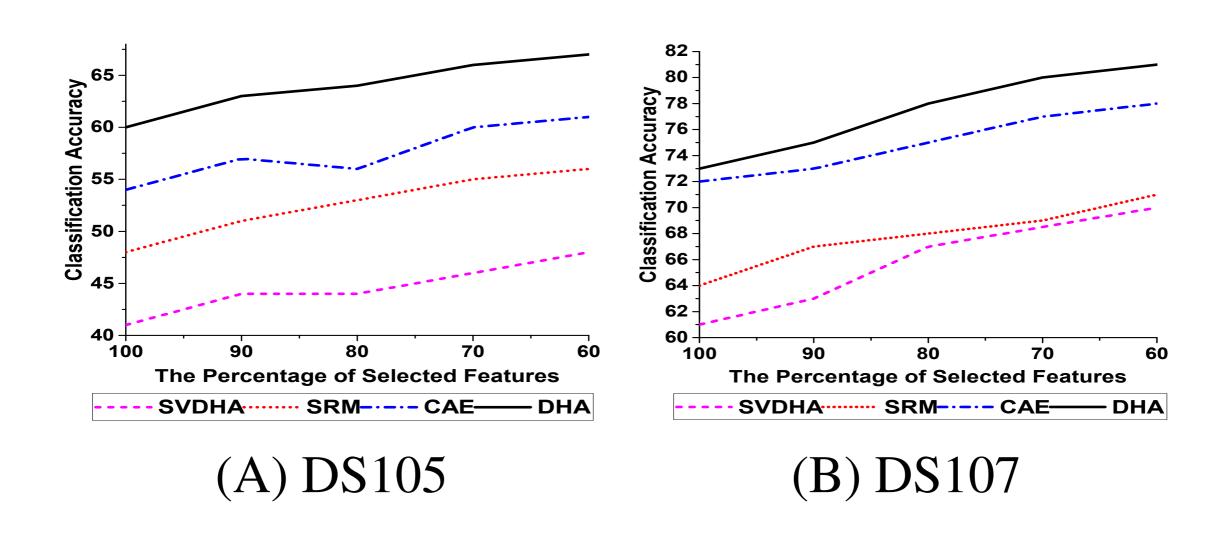


Figure 1: Comparison of different HA algorithms on complex task datasets by using ranked voxels.

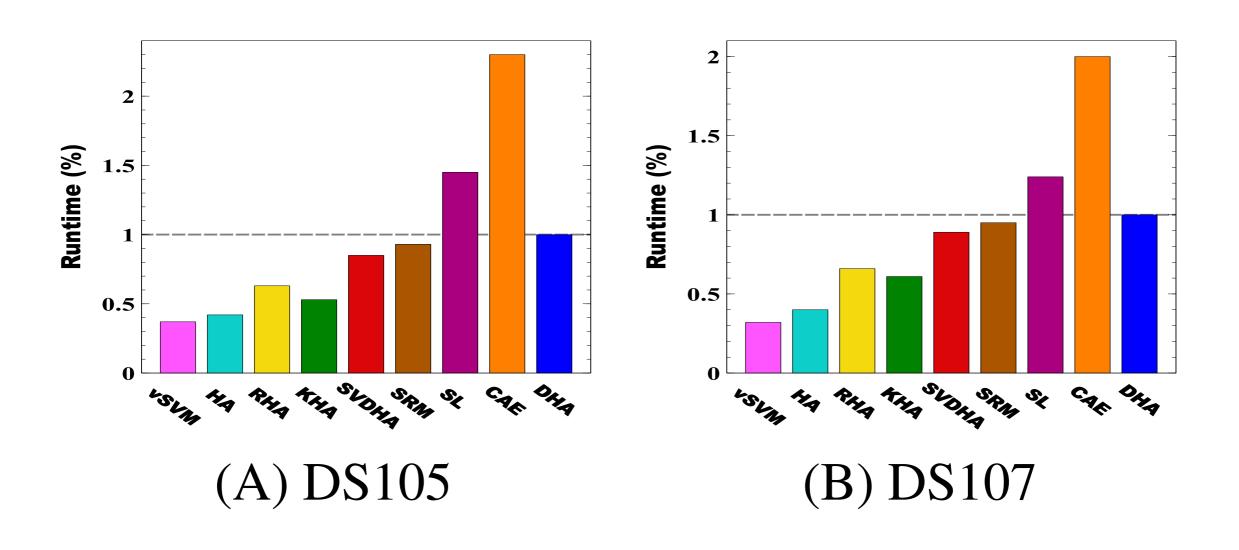
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Classification analysis by using feature selection



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Runtime Analysis



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Future Works

- **★** This paper extended a deep approach for hyperalignment methods in order to provide accurate functional alignment in multi-subject fMRI analysis.
- ★ Deep Hyperalignment (DHA) can handle fMRI datasets with nonlinearity, high-dimensionality (broad ROI), and a large number of subjects. Further, its time complexity fairly scales with data size and the training data is not referenced when DHA computes the functional alignment for a new subject.
- **★ In the future, we will plan to employ DHA for improving the performance of other techniques in fMRI analysis, e.g. Representational Similarity Analysis (RSA).**

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Thank You!

Q & A

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