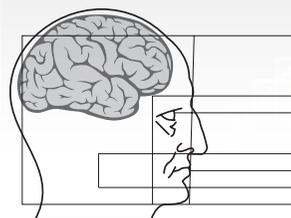
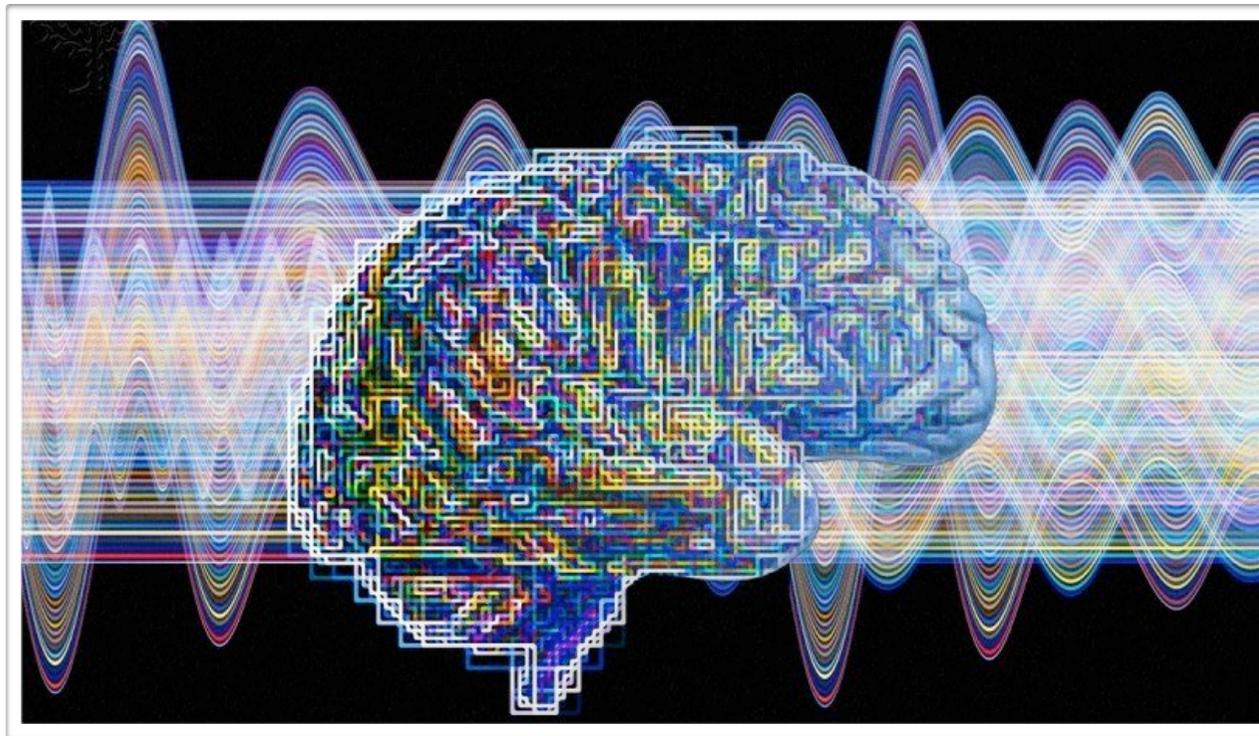


The multivariate partial least squares (PLS) framework for neuroimaging



FU Berlin, Sept 1, 2015



In search of multivariate data “patterns” in neuroimaging

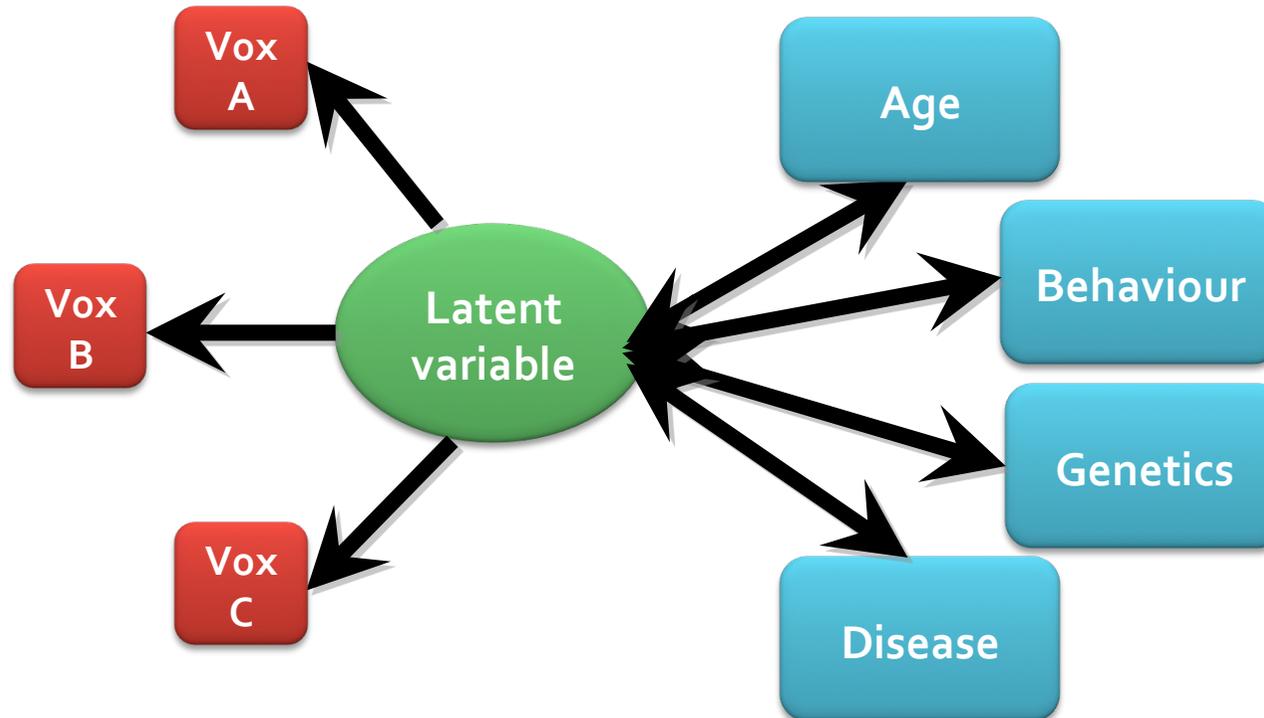
- Multivariate techniques make using complex imaging data simpler.
 - Leverage vast information to maximize our understanding of phenomena of interest.
- MVPA:
 - Often purported in neuroimaging as a multivariate method, but is often utilized as a univariate technique (e.g., linear discriminant and logistic models are typically *dimensionally* univariate).
- What about truly multivariate (multidimensional) models?

Partial least squares (PLS)

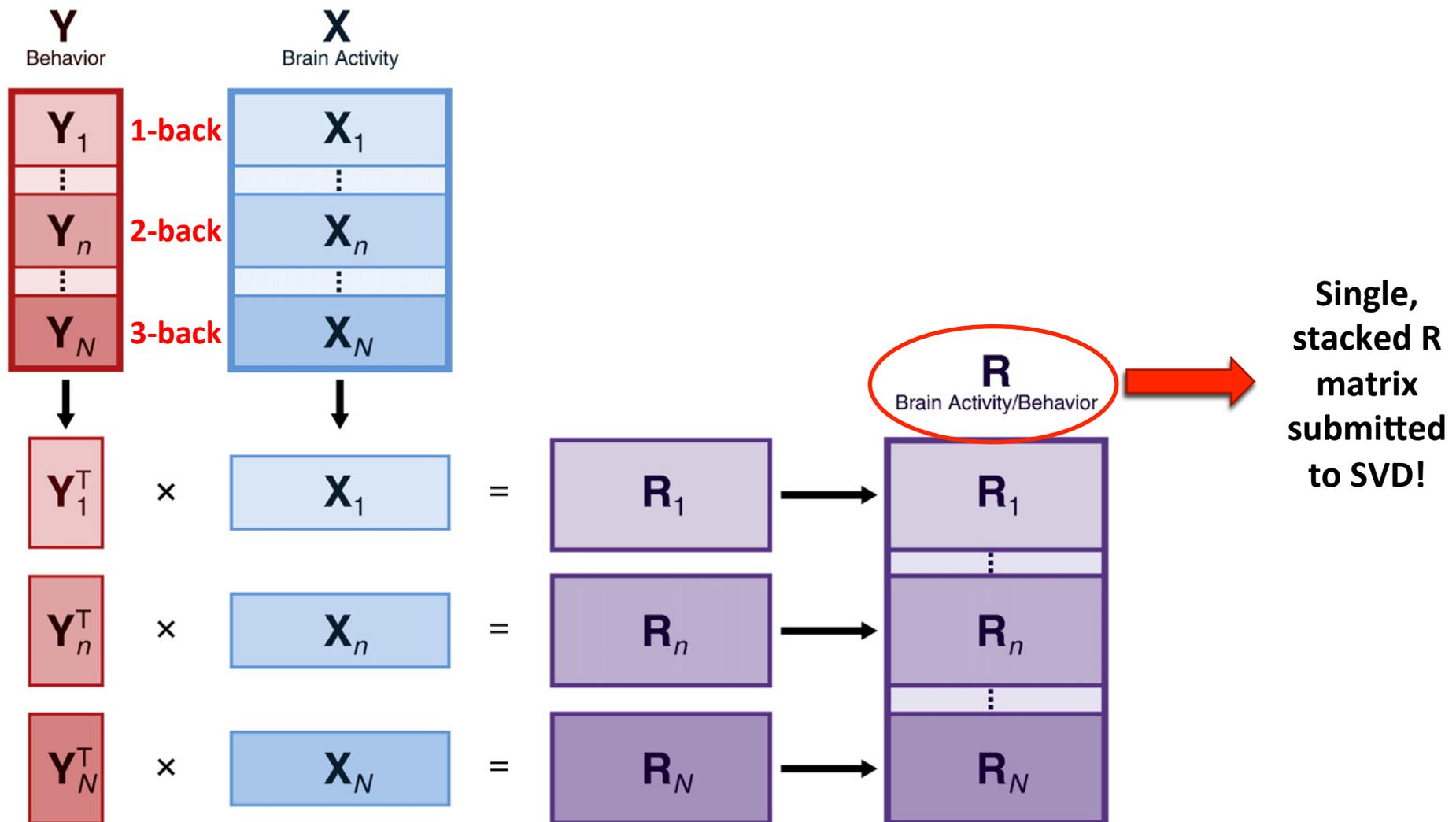
- Is a general multivariate (multidimensional) statistical method:
 - Whereas MVPA often leverages multiple sources of brain activity to discriminate between discrete classes/groups/states;
 - PLS (McIntosh et al., 1996) is more general in form, allowing researchers to find multivariate, latent-level “patterns” linking brain data to *any other* variables of interest (classes, continuous variables, etc.) in one mathematical step.
 - Can be utilized in the context of EEG, fMRI (block design, event-related), structural MR, PET, network indices, etc.

PLS: What's to gain?

- PLS is an effective way of **reducing dimensionality**
- Moves us to **latent space**: allows us to capture very complex phenomenon in fewer dimensions than univariate = PARSIMONY



PLS: Data set up...



The SVD

- ➔ SVD (like PCA, but for rectangular matrices) then produces orthogonal latent variables that optimally express relations between X and Y.
- ➔ Singular values rank-ordered by strength.

$$\mathbf{SVD\ of\ } R_{XY} = \mathbf{U\ S\ V'}$$

Condition/group/
class/behaviour
weights

Singular
values

Voxel
weights

SVD: How many dimensions are mathematically possible?

- ➔ # is always equal to the **smaller** rank of X (e.g., voxel measures) or Y (e.g., behavioural measures/conditions)
- ➔ Ask the audience! How many dimensions possible here (ignoring subjects)?
- ➔ 6 (2 behav measures*3 conditions; brain=12 vox*3 conditions)

		Voxels												
$\mathbf{X} = \begin{bmatrix} \mathbf{X}_1 \\ \mathbf{X}_2 \\ \mathbf{X}_3 \end{bmatrix}$	s1	2	5	6	1	9	1	7	6	2	1	7	3	1-back
	s2	4	1	5	8	8	7	2	8	6	4	8	2	
	s3	5	8	7	3	7	1	7	4	5	1	4	3	
	s1	3	3	7	6	1	1	10	2	2	1	7	4	2-back
	s2	2	3	8	7	1	6	9	1	8	8	1	6	
	s3	1	7	3	1	1	3	1	8	1	3	9	5	
	s1	9	0	7	1	8	7	4	2	3	6	2	7	3-back
	s2	8	0	6	5	9	7	4	4	2	10	3	8	
	s3	7	7	4	5	7	6	7	6	5	4	8	8	

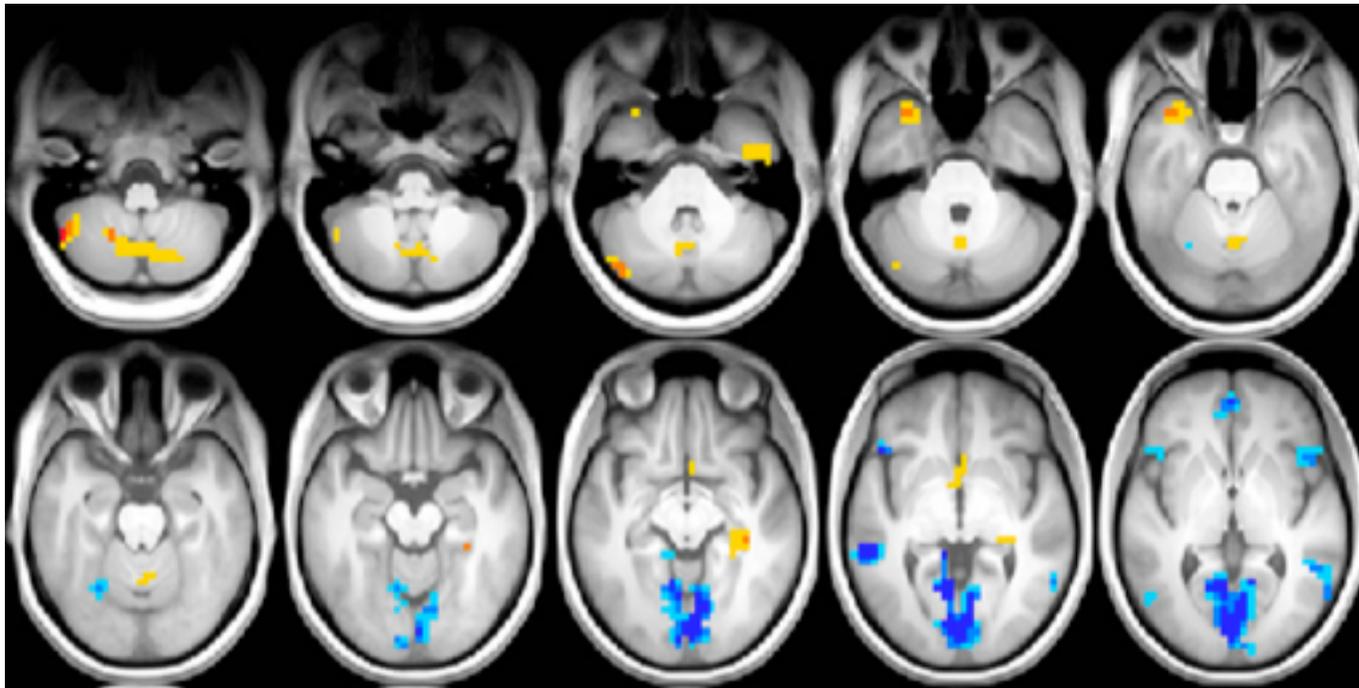
		ISD _{RT} Mean _{RT}		
$\mathbf{Y}_{\text{behavior}} = \begin{bmatrix} \mathbf{Y}_{\text{behavior},1} \\ \mathbf{Y}_{\text{behavior},2} \\ \mathbf{Y}_{\text{behavior},3} \end{bmatrix}$	s1	15	600	1-back
	s2	19	520	
	s3	18	545	
	s1	22	426	2-back
	s2	21	404	
	s3	23	411	
	s1	29	326	3-back
	s2	30	309	
	s3	30	303	

We've gained parsimony, but what else?

- Multiple comparisons problem is minimized.
- “Brute force” of univariate approach not required.
- As in MVPA, brain data are leveraged together, but PLS simply extends this logic to an arguably more flexible model class than typically employed in the MVPA world.

PLS: Another bonus...

- Brain weights can be either positive or negative across voxels within a single dimension
- Unlike many univariate designs/packages, can visualize both positive and negative voxel effects simultaneously.

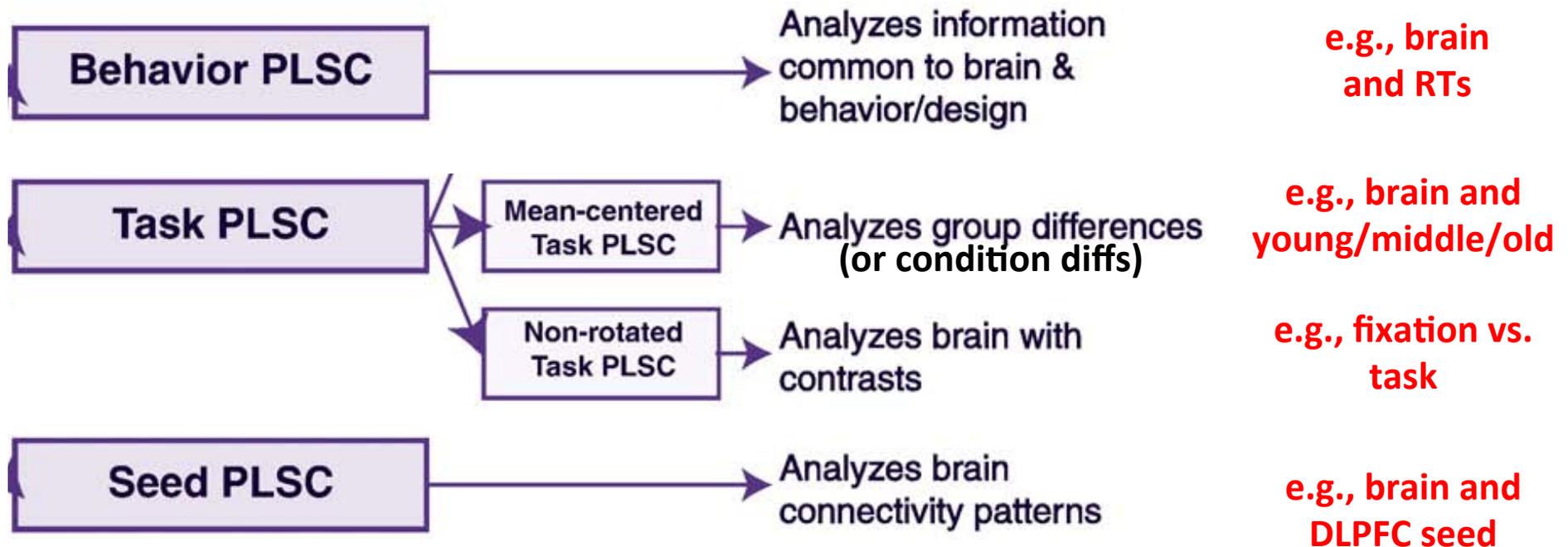




PLS: Practical details...

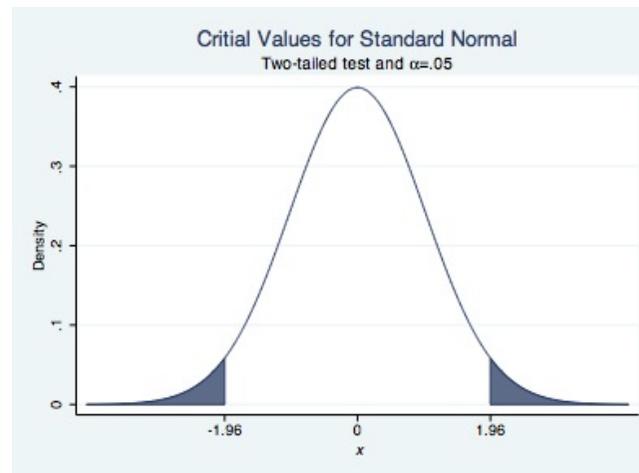


Typical forms of PLS



Model evaluation

- SVD as described above = **fixed effect model**; so how can we generalize to the population (like random effects models)?
- Inferential analytical approaches are available, but arguably make too many parametric assumptions to be used routinely across broad types of data. Why assume shape when we can evaluate using **nonparametric** methods?



Two-stage nonparametric model evaluation

1. Dimensions to keep? Permutation on singular values

- A singular value represents the “strength” (like variance accounted for in the data) of a particular dimension of XY relations; how strong is strong enough?
- We can test the robust strength of each singular value by randomly shuffling rows of X without changing Y, and test whether singular value would be just as high.
- Run new SVD on this reshuffled dataset, get singular value; do that 1000 times to get distribution of singular values
 - If get singular value as strong as in original data $< .05$, then keep that dimension!

At voxel level...

- Now that we have chosen which latent variables to keep, we need to “threshold” which brain data (e.g., voxels) to reliably report within those latent variables.
- 2. Stage 2 then = **Bootstrapping** (with replacement)
 - Reach into sample hat, pull out subject x; put back in, draw again, etc., until have a new “resample” of same size as original.
 - Do that 1000 times, running SVD on each resample
 - Derive bootstrap standard errors for each voxel.
 - Original voxel weight/bootstrap standard error = **BOOTSTRAP RATIO (BSR)**. Tells us how probable it would be that this voxel is “really” active, across multiple resamples.
 - Typical choice: BSR=3.00 (akin to z-score).

PLS: Individual differences

➤ Latent variables chosen, voxels thresholded...
but what about the subject level?

➤ “Brain score”

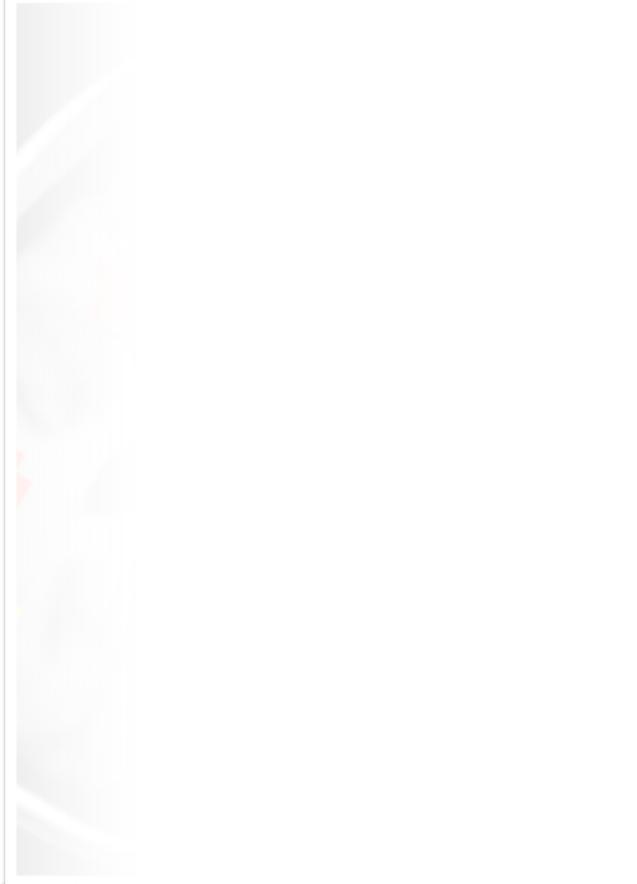
$$\text{SVD of } R_{XY} = U S V'$$

Condition/
group
weights

Singular
values

Voxel
weights

$$\text{Brain Score}_i = V_j * \text{Voxel Data}'_{ij}$$

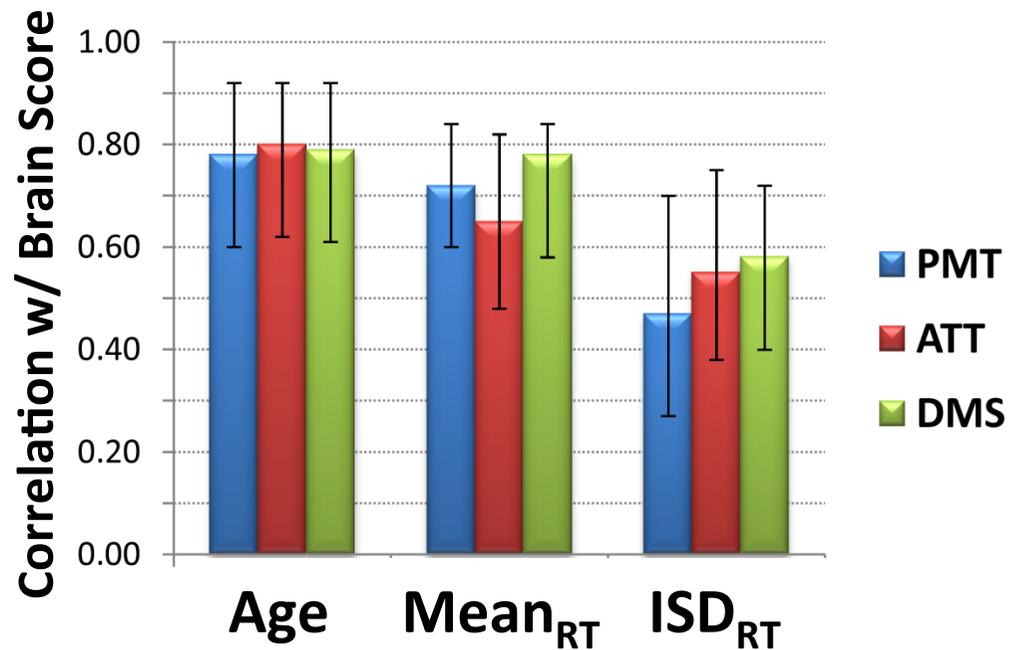
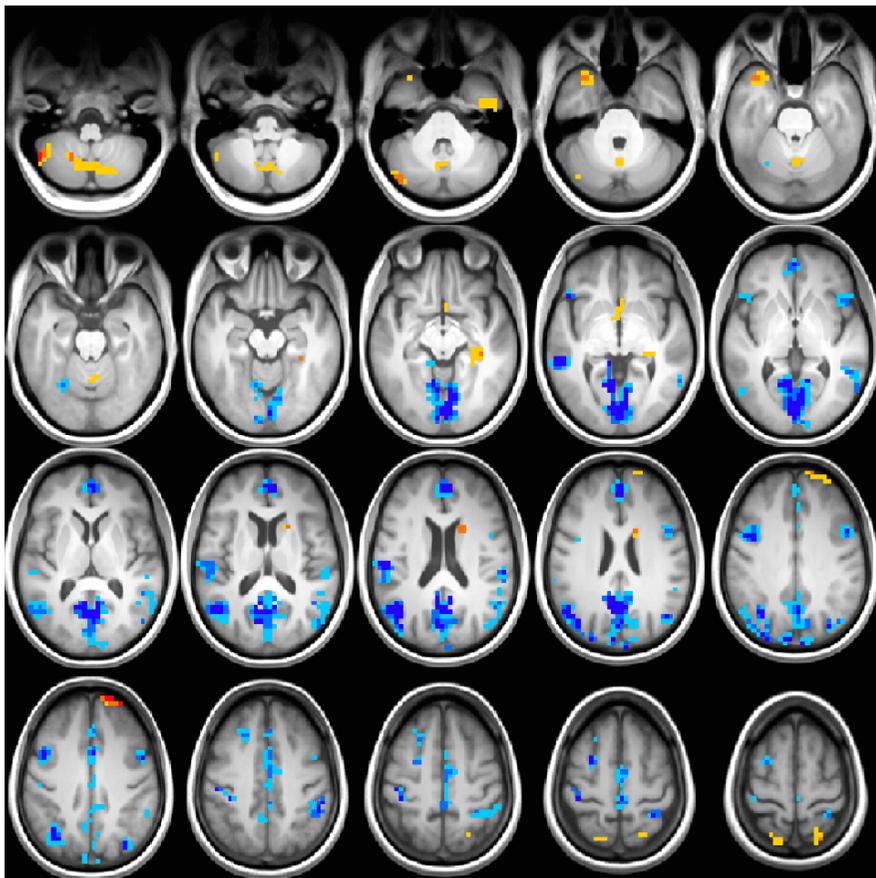


PLS: Published examples



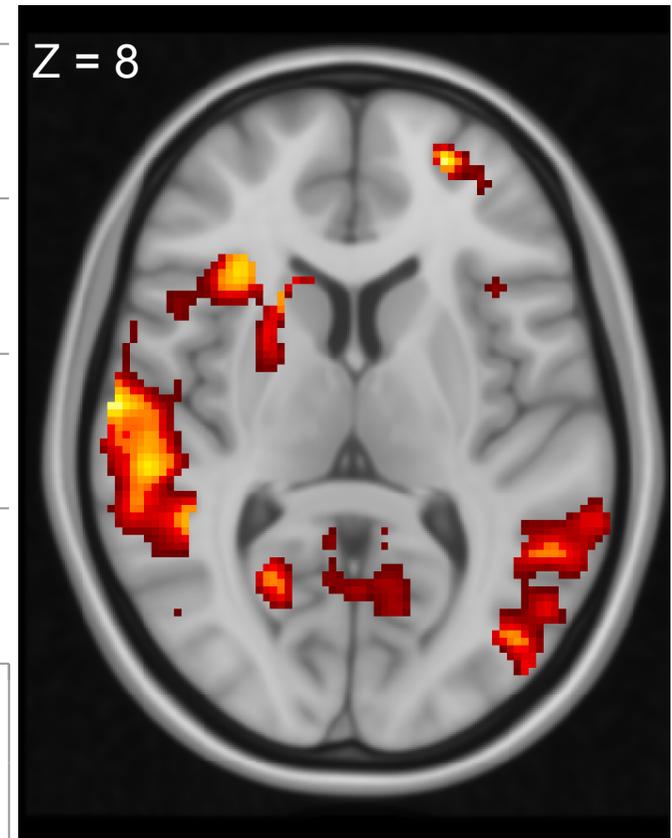
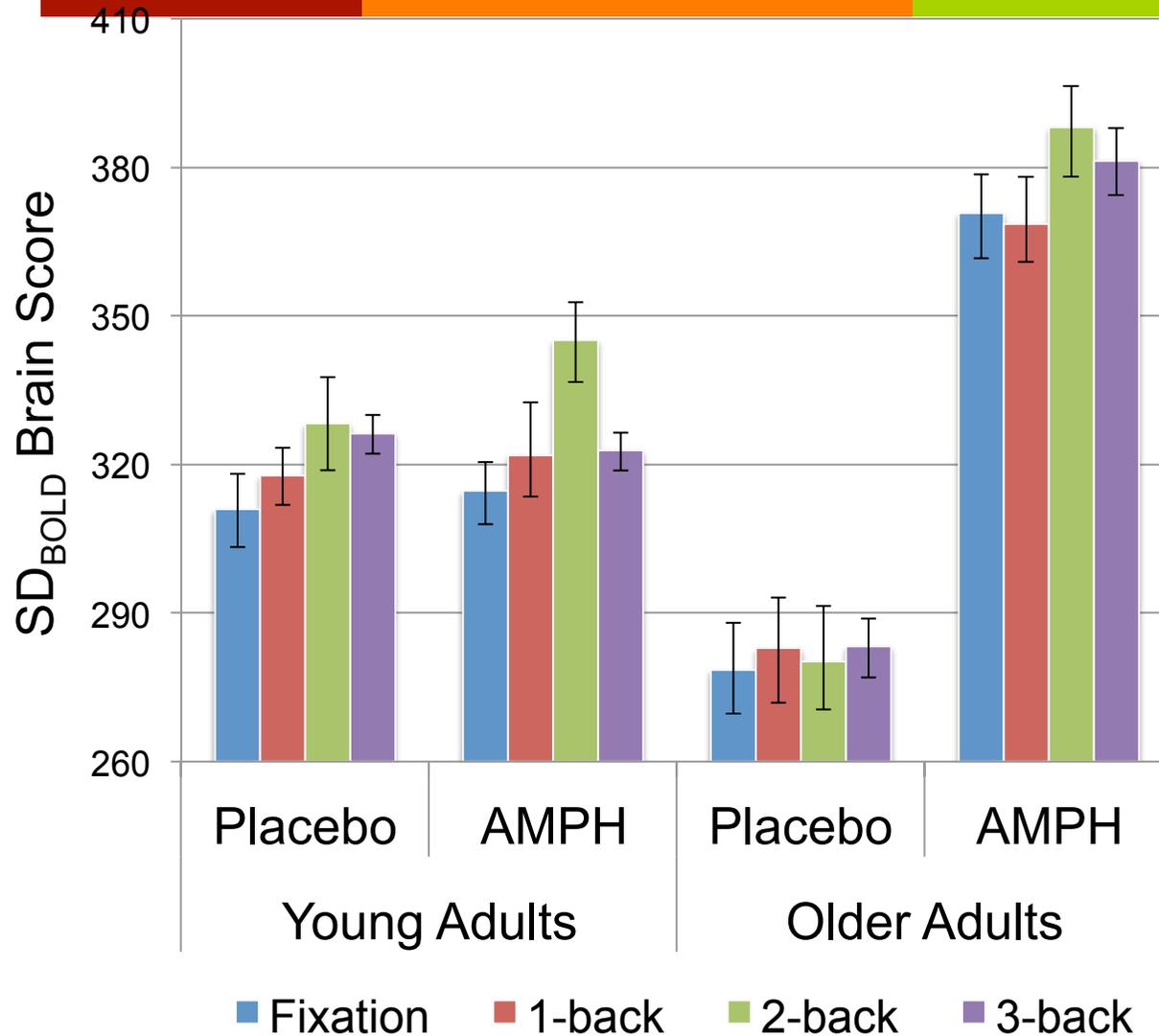
Example #1: Linking BOLD signal variability to age, RT_{mean} , RT_{sd} on three cognitive tasks

- Young, fast, stable adult performers = higher SD_{BOLD} (single robust LV).



Garrett et al. (2011), *JNeurosci*

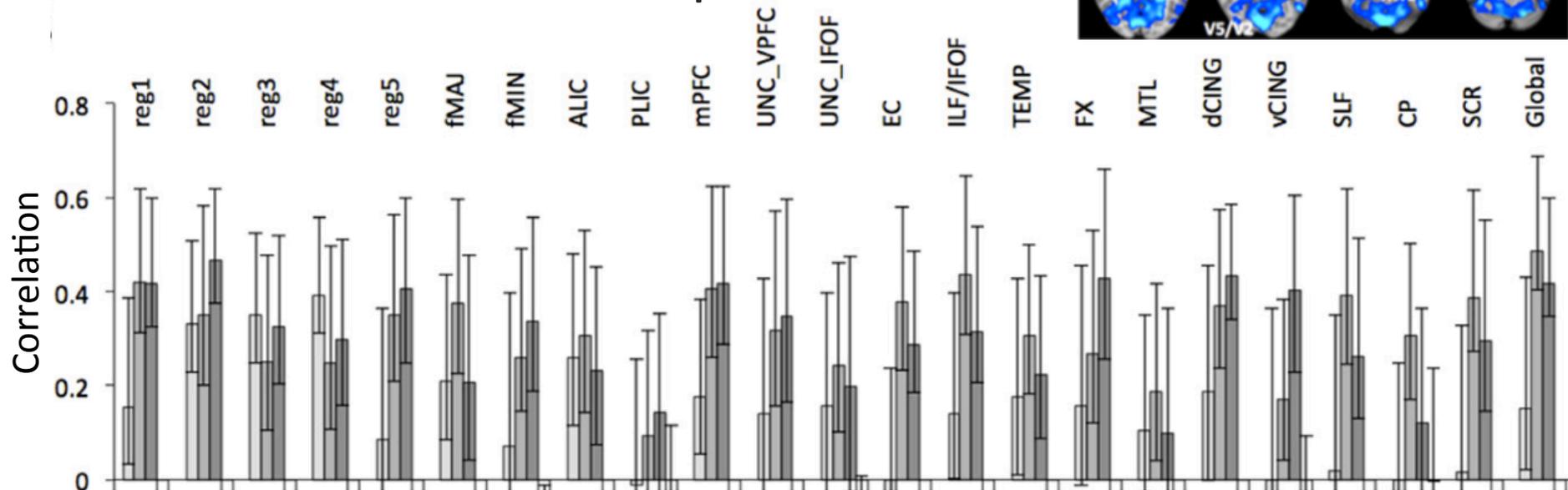
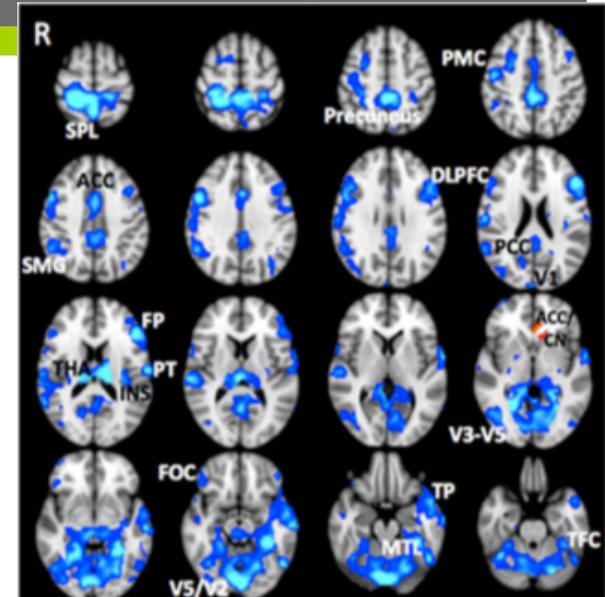
Example #2: Can AMPH boost SD_{BOLD} in older adults? (Garrett et al., 2015, *PNAS*)



Example #3: Linking FA and mean_{BOLD}

(*Burzynska, *Garrett, et al., 2013, *JNeurosci*)

- Single robust LV; Higher FA correlated with lower BOLD during n-back (1-, 2-, 3-back)
- Higher FA, lower n-back BOLD, better behavioral performance.



Summary

- PLS is a useful technique for reducing data dimensionality and finding “patterns,” and can capture broad scale relations between any brain data and any other variables of interest.
- Given non-parametric assumptions, there are no real bounds on the types of variables that can be examined.
- Can be either exploratory or hypothesis driven; it does what you tell it to do!

Many thanks to all!

➤ McIntosh PLS software: Google “PLS Rotman...” and the latest release will come up.

➤ Primary intro references:

1. Krishnan, A., Williams, L. J., McIntosh, A. R., & Abdi, H. (2011). Partial Least Squares (PLS) methods for neuroimaging: a tutorial and review. *NeuroImage*, *56*(2), 455–475.
2. McIntosh, A. R., & Lobaugh, N. J. (2004). Partial least squares analysis of neuroimaging data: applications and advances. *NeuroImage*, *23 Suppl 1*, S250–63.
3. McIntosh, A. R., Bookstein, F. L., Haxby, J. V., & Grady, C. L. (1996). Spatial pattern analysis of functional brain images using partial least squares. *NeuroImage*, *3*(3 Pt 1), 143–157.
4. McIntosh, A. R., Chau, W. K., & Protzner, A. B. (2004). Spatiotemporal analysis of event-related fMRI data using partial least squares. *NeuroImage*, *23*(2), 764–775.