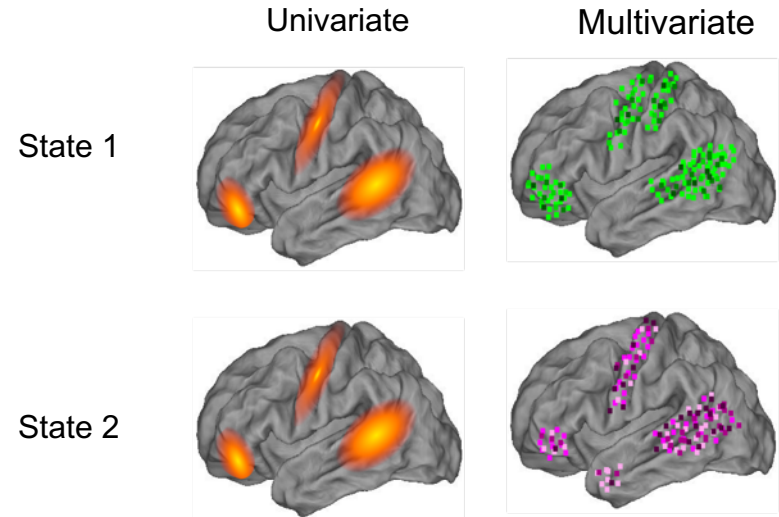
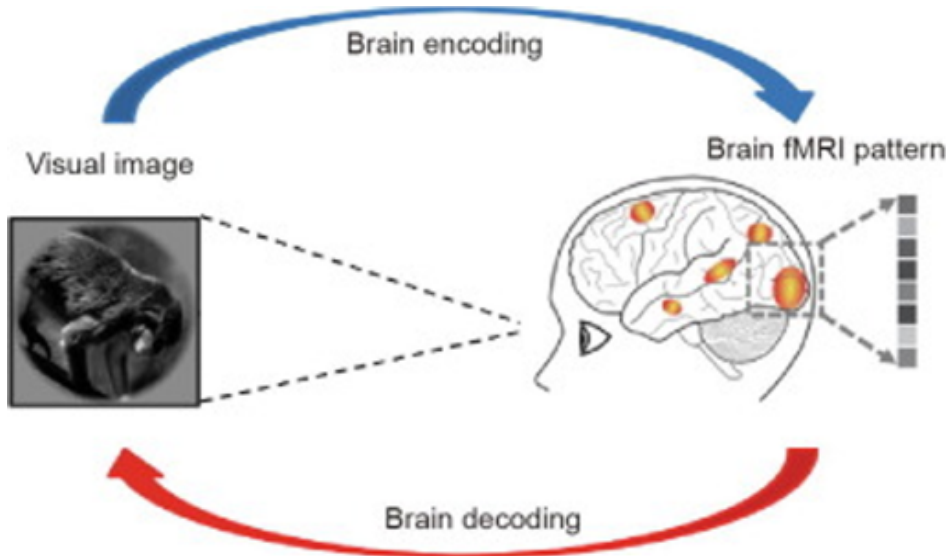


Statistical Pattern Recognition with fMRI

Heini Saarimäki
Postdoctoral Fellow
Human Information Processing Laboratory
Tampere University

Multivariate / multivoxel pattern analysis (MVPA), Pattern recognition, Machine learning, Decoding, Classification, ...



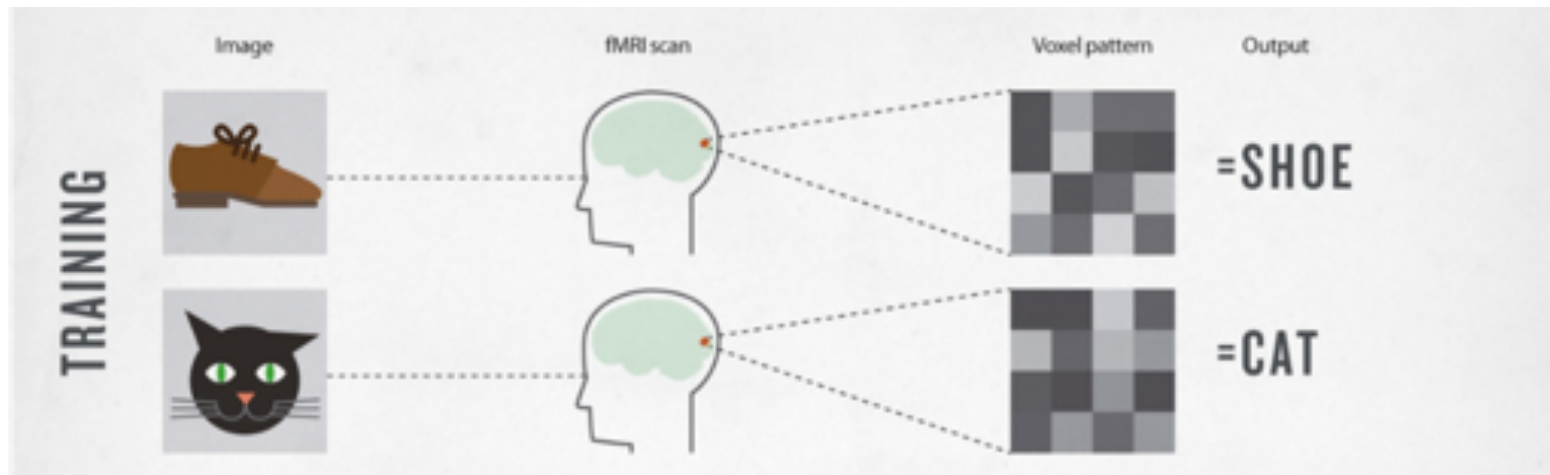
Outline

WHY: When and why to use classification methods

WHAT: Nuts and bolts of pattern classification

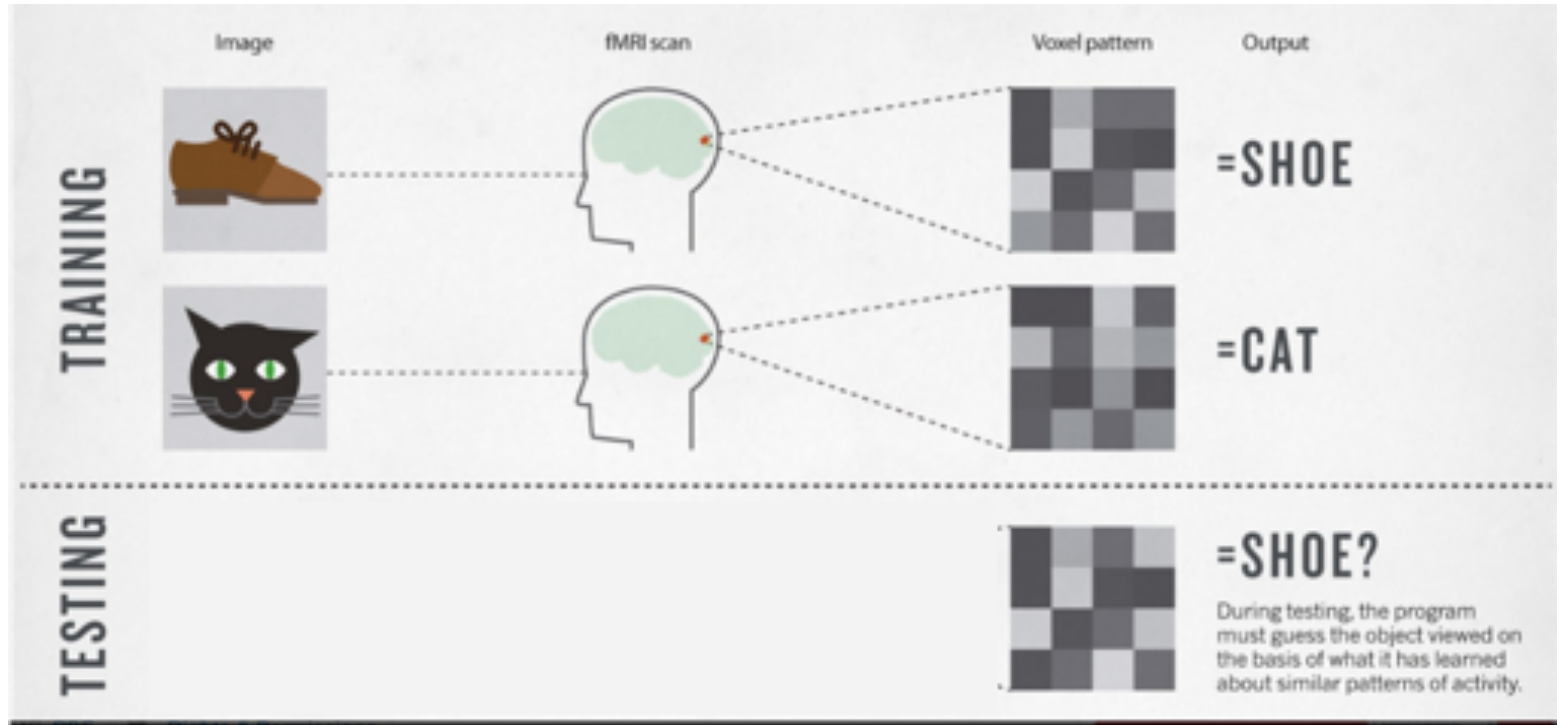
HOW: Overview of the workflow

Simplified framework:



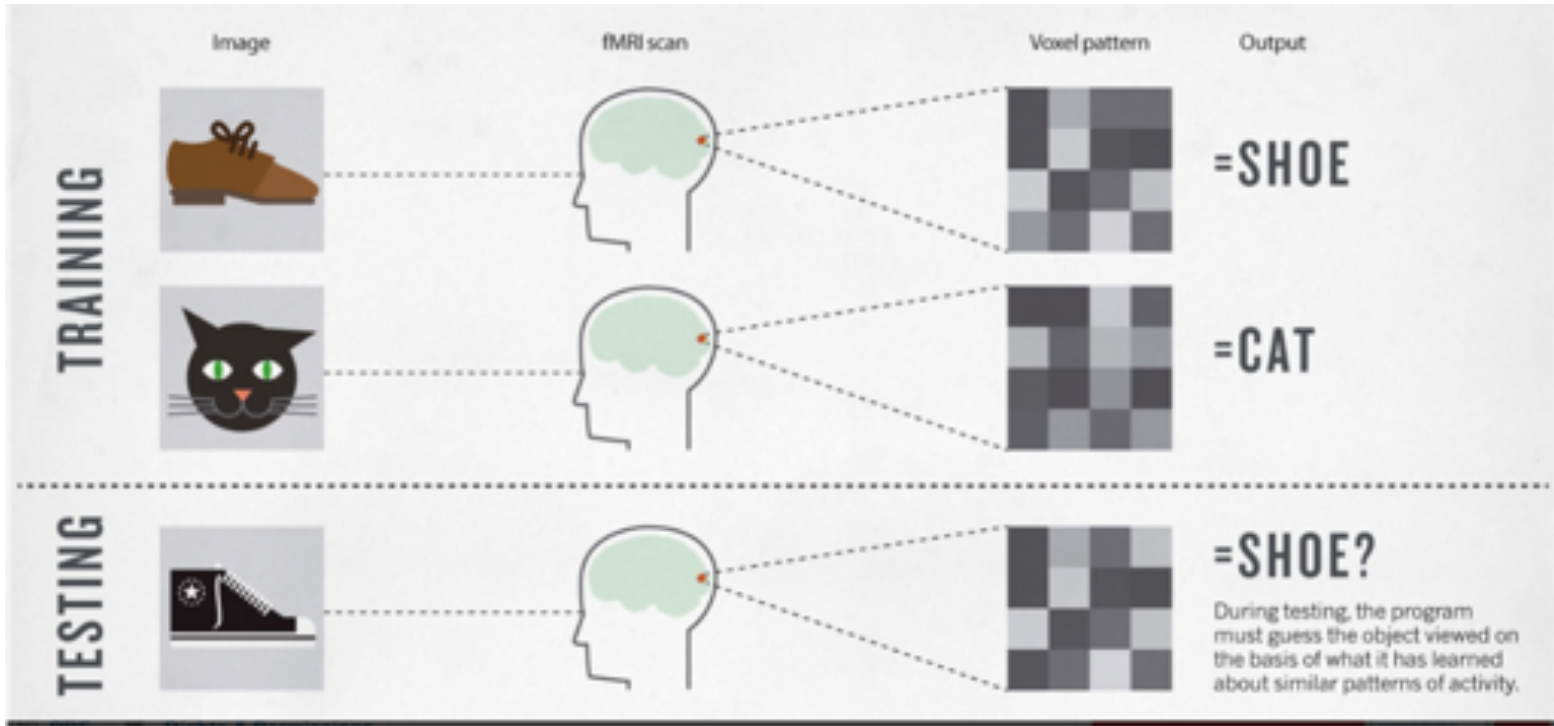
e.g. Haxby et al. (2001) *Science*
Figure credits: Smith (2013) *Nature*

Simplified framework:



e.g. Haxby et al. (2001) *Science*
Figure credits: Smith (2013) *Nature*

Simplified framework:

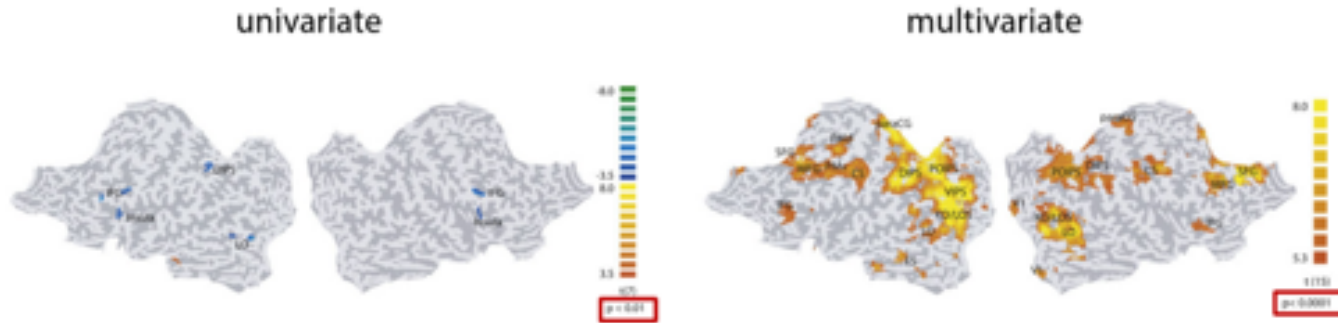


e.g. Haxby et al. (2001) *Science*
Figure credits: Smith (2013) *Nature*

Why pattern recognition?

Higher sensitivity when compared to univariate analyses → detects pattern differences when activation overlaps

Example: Representation of perceptual choices

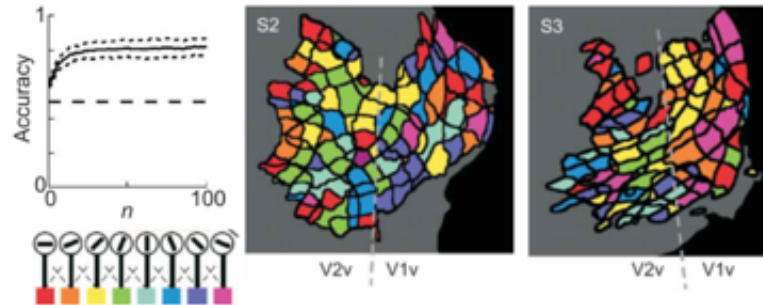


Why pattern recognition?

Higher sensitivity when compared to univariate analyses → pattern differences when activation overlaps

Can study representational content in a brain region rather than general activation (also activity vs information)

Example: Representation of orientations



Why pattern recognition?

Higher sensitivity when compared to univariate analyses → pattern differences when activation overlaps

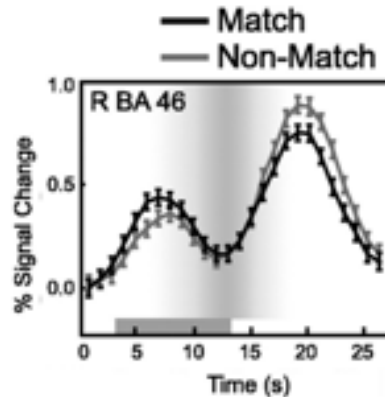
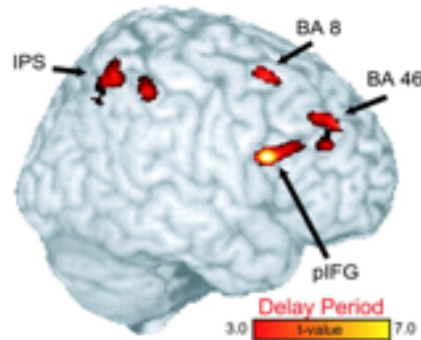
Can study representational content in a brain region rather than general activation (also activity vs information)

Methods readily available from machine learning

The principle is relatively easy to understand

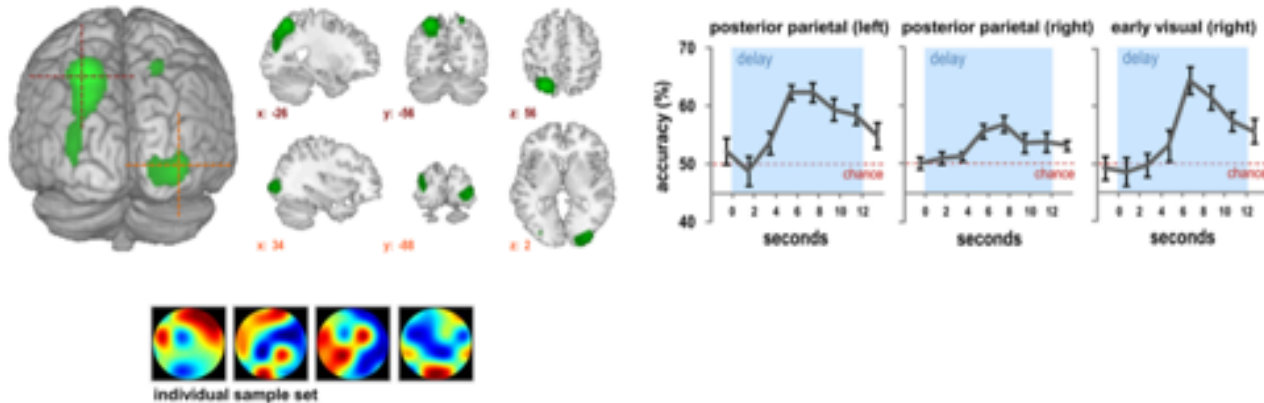
Activity vs information

Activity: Tells us about general involvement in cognitive function (e.g. working memory)



Activity vs information

Information: Tells us about representational content (e.g. memory trace of A vs. memory trace of B)



Applications

Task classification, testing theories of representational structure

Medical diagnosis classification

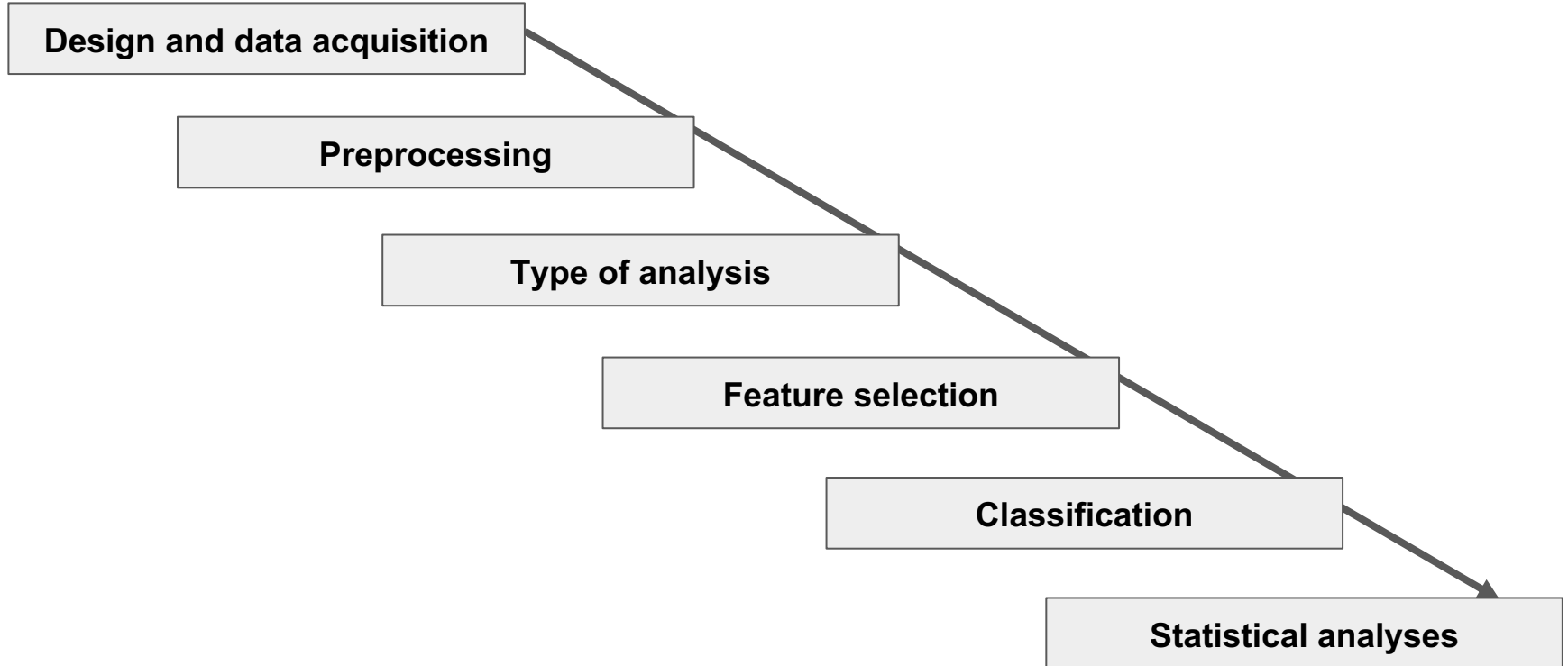
Medical Diagnosis

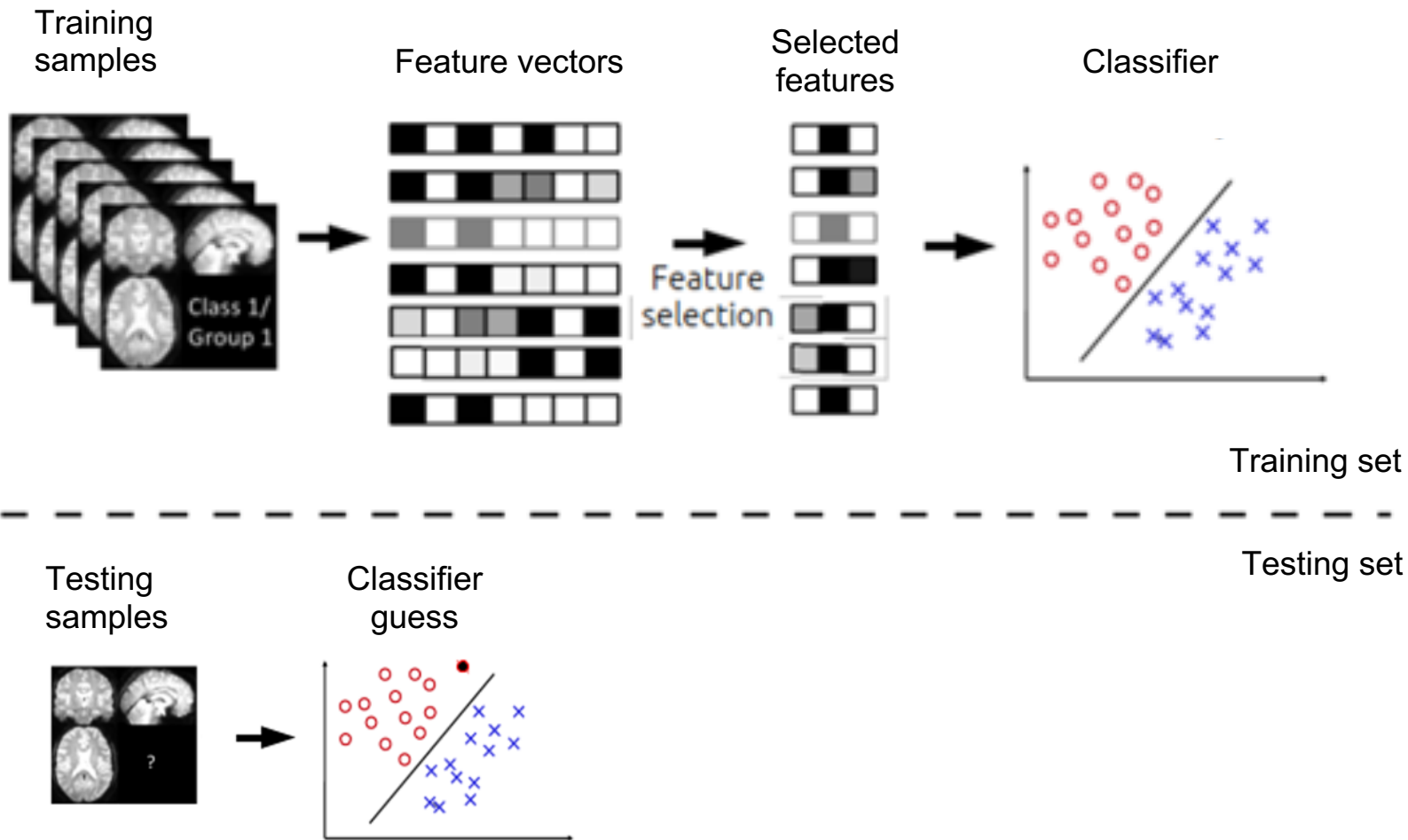


Brain-Computer-Interface



Overview of the workflow





Step 1: Design and data acquisition

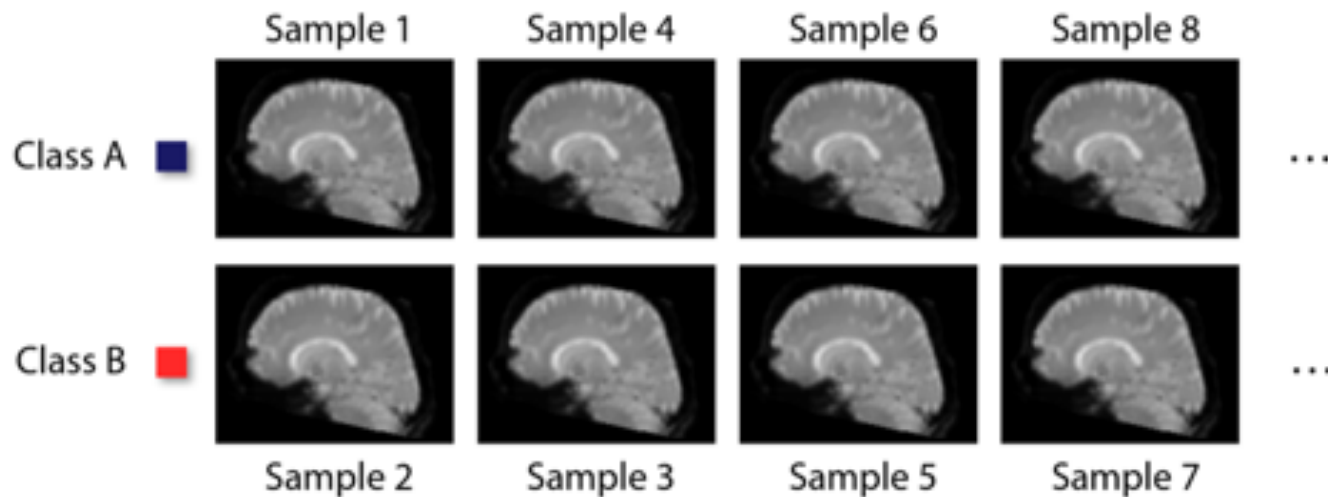
Categories



Key concepts: Sample

Samples are data that belong to a class

Examples: EPI volumes, beta volumes, VBM maps, EEG data

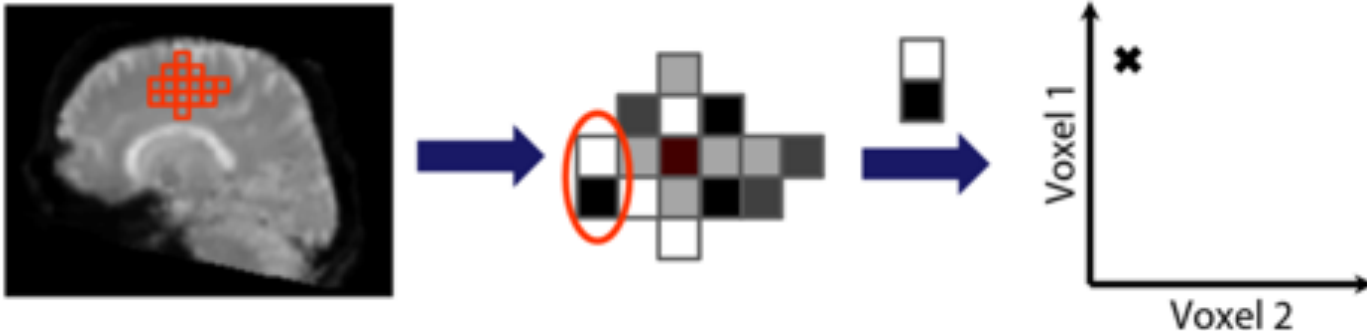


Key concepts: Feature

Each feature is a measured variables that can be used for classification

- Each feature (hopefully) aids the classification process, by contributing signal and/or suppressing noise
- Each feature spans up a dimension \rightarrow they build the feature space

Examples: A voxel, connectivity graph, EEG channel



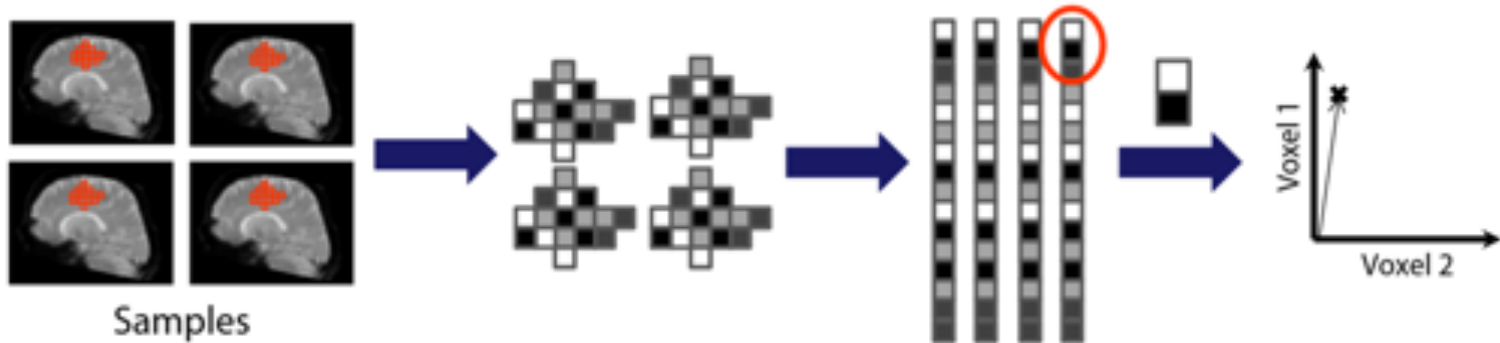
Key concepts: Pattern

A pattern is a sample for a set of features

A pattern is a point (or vector) in p -dimensional space (p is # of features)

Alternative uses of term "pattern" with different meaning:

- Prototypical pattern (i.e. the true class mean)
- Discriminating pattern (function that discriminates classes)



Key concepts: Label

A label denotes the class membership of a pattern with a number

For classification the number is categorical and often arbitrary (some classifiers require 0 and 1 or -1 and 1)

For regression the number denotes a continuous number which is the regression target

Class A		Class B	
Label: 1		Label: -1	

Step 2: Preprocessing

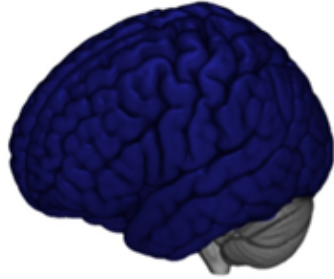
Minimal preprocessing

No smoothing

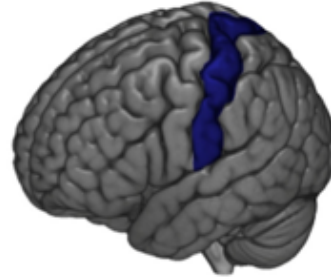
Normalization of signal intensities

Step 3: Type of analysis

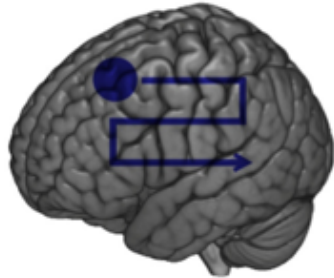
Wholebrain



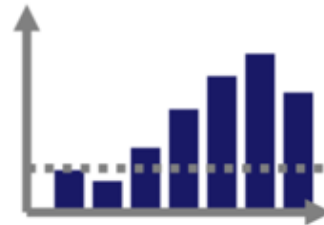
Region of Interest



Searchlight



time-resolved

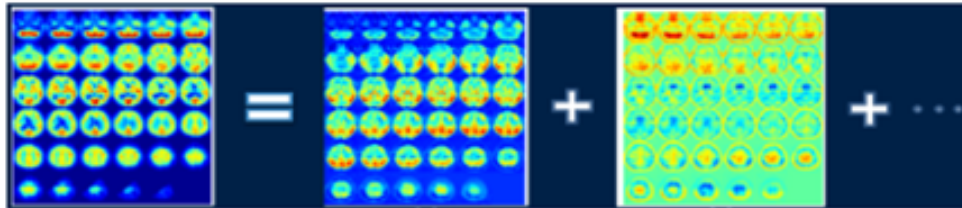


Step 4: Feature selection

ANOVA

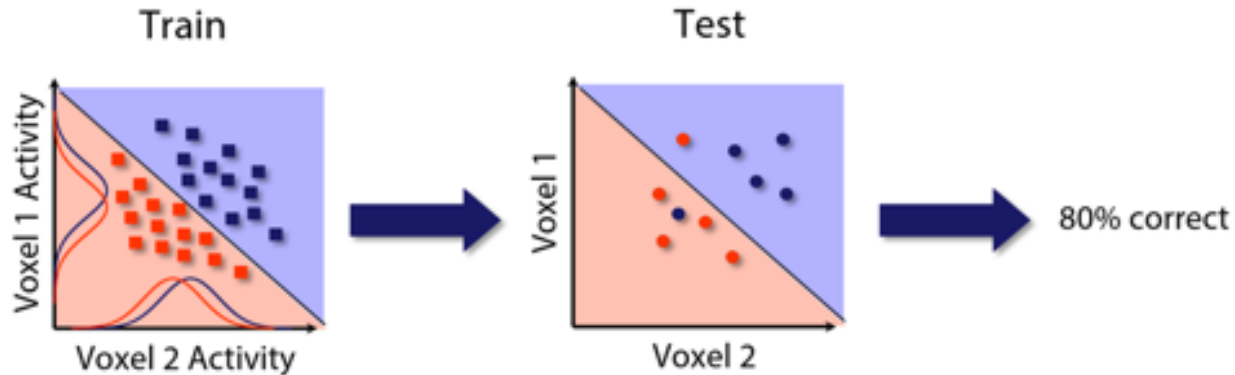


PCA



Step 5: Classification

1. Separate data in training and test data
2. "Train" a classifier (i.e. find a good separating line)
3. Apply this classifier (i.e. the line) to test data
4. Result: Accuracy of test data (i.e. % correct prediction)



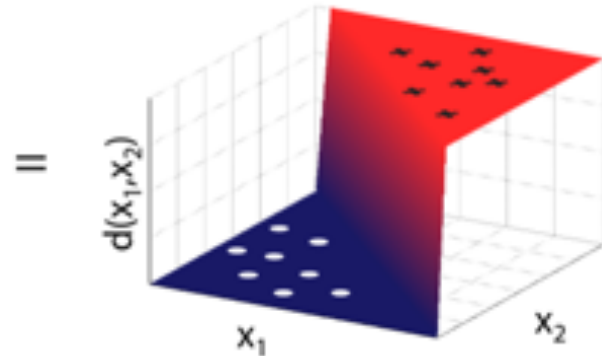
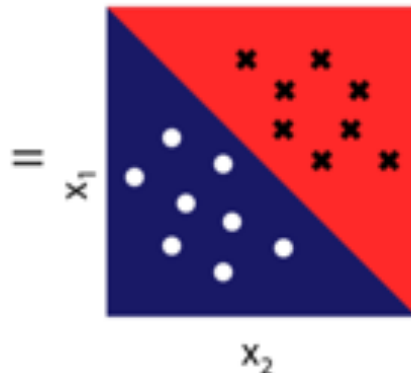
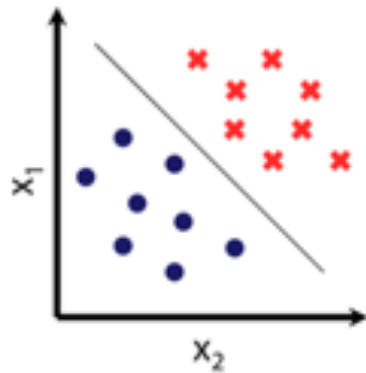
Key concepts: Classifier

A function that separates feature space

Example for one sample with two features: $f(x_1, x_2) = -0.5$

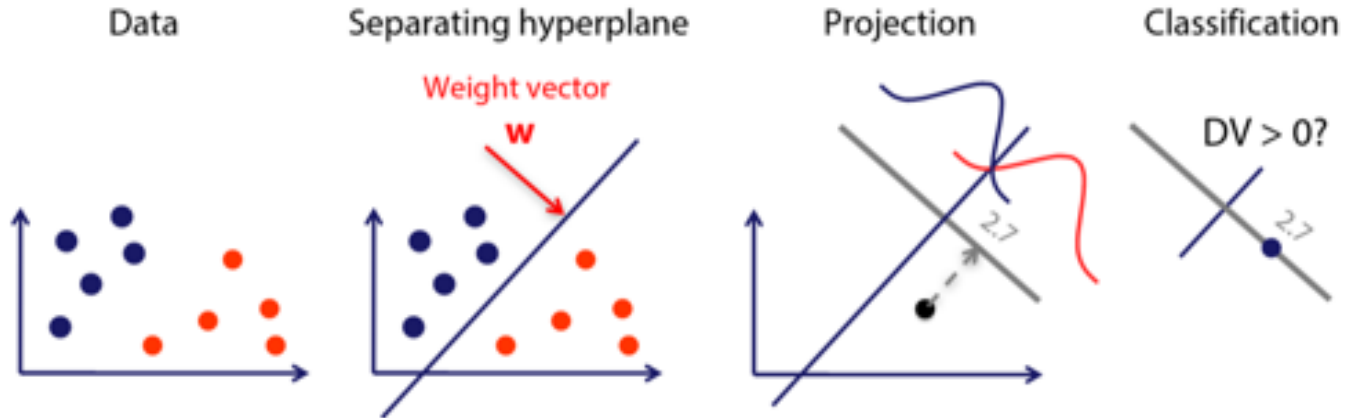
This decision value f is then binarized in a decision function:

if $f(x_1, x_2) > 0$: $d(x_1, x_2) = \mathbf{1}$; *if* $f(x_1, x_2) \leq 0$: $d(x_1, x_2) = \mathbf{-1}$



Step 5: Classification

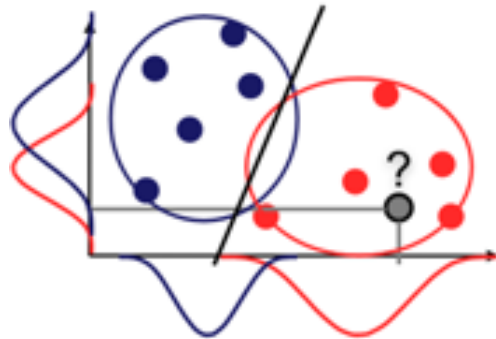
Geometric intuition



$$f(x) = w^T x + b$$

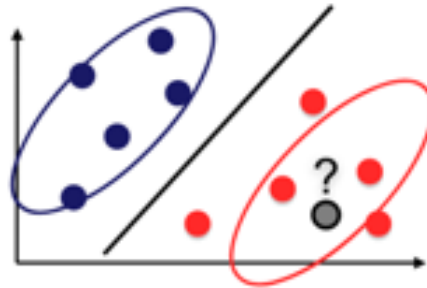
Step 5: Classification - linear classifiers

Gaussian Naïve Bayes



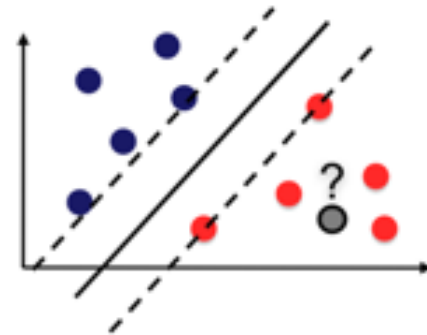
Ignores covariance between voxels

Linear Discriminant Analysis



Considers covariance between voxels

Support Vector Machine

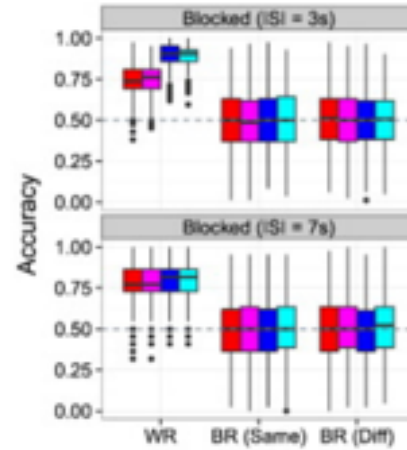
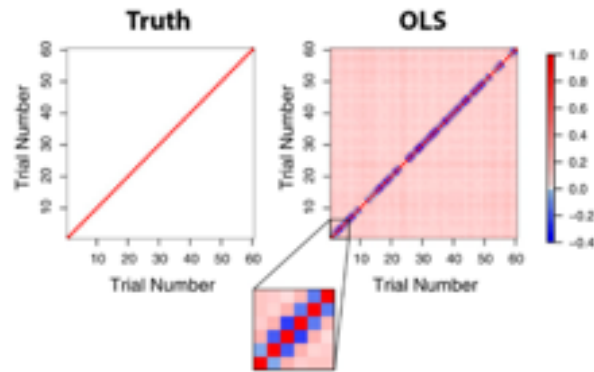


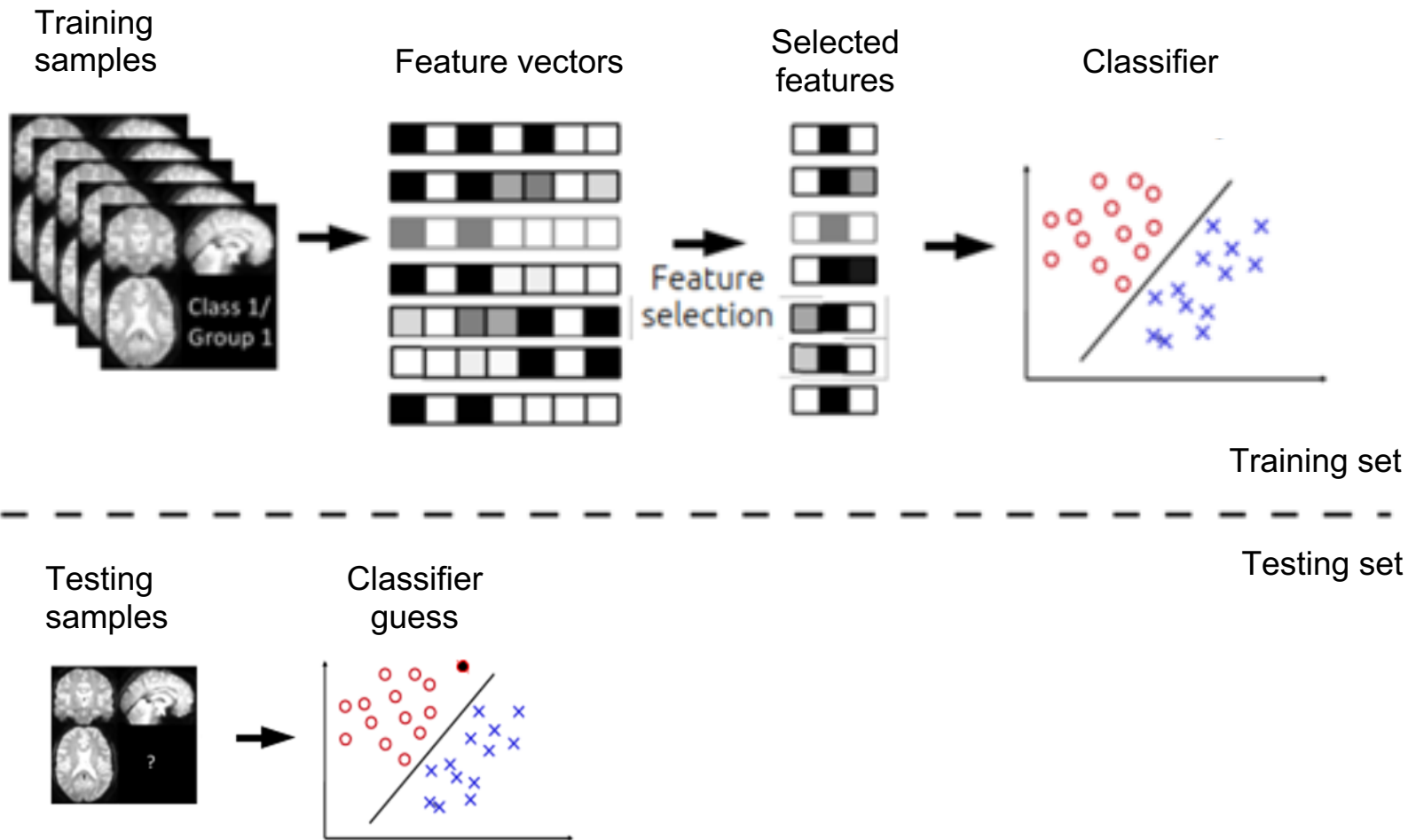
Maximizes margin (distance between closest points of different classes)

Step 5: Classification - cross-validation

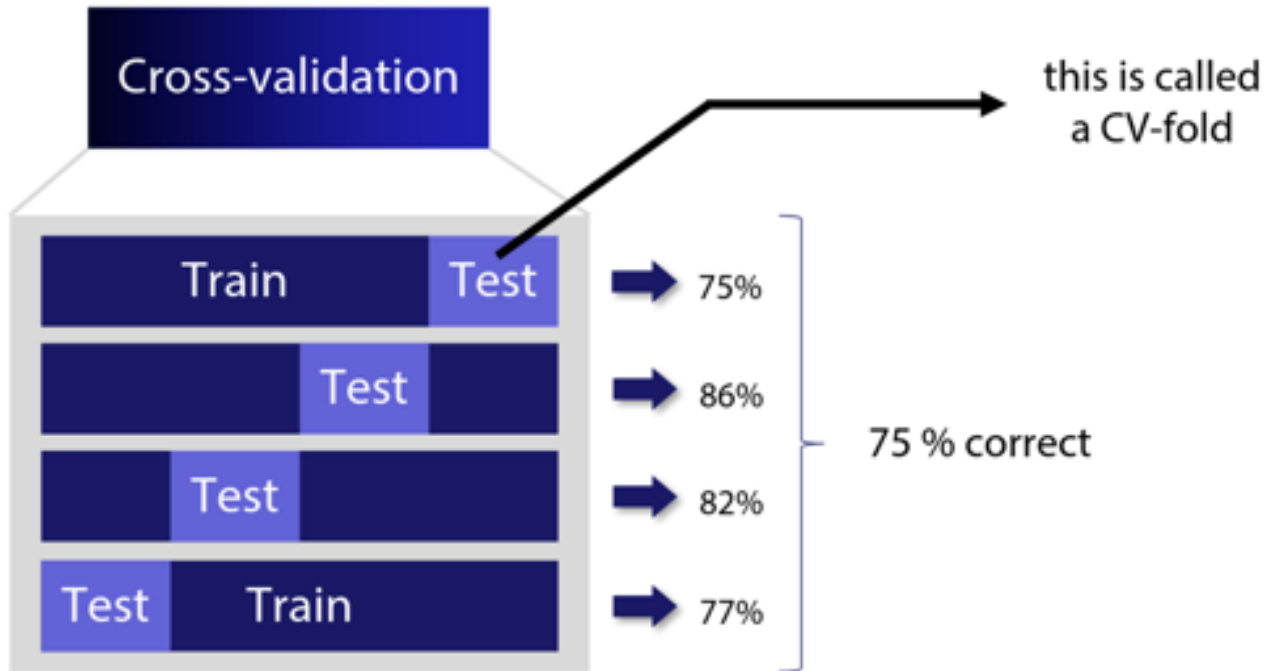
Typical Analysis: Leave-one-run-out cross-validation

Reason: Non-independence within run can bias results





Step 5: Classification - cross-validation



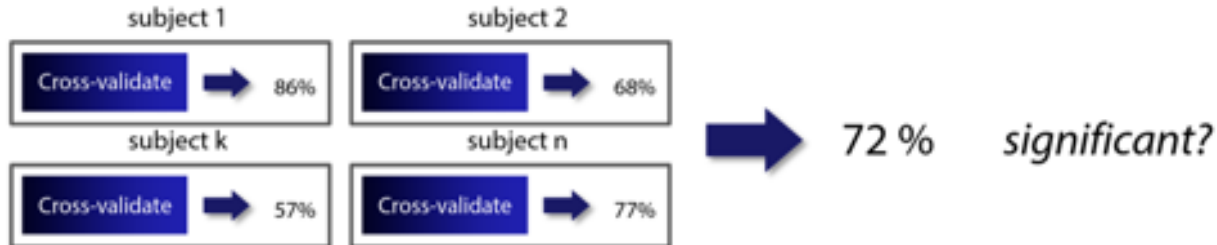
Step 6: Statistics - classifier accuracy

“Decoding level”



- within-subject classification (condition A vs. condition B)
- between-subject classification (group A vs. group B)

“Second level” (group analysis)



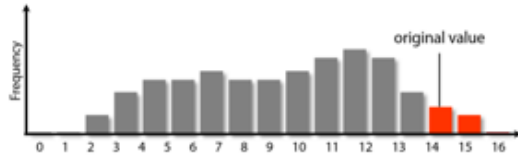
Step 6: Statistics - permutation tests

Procedure:

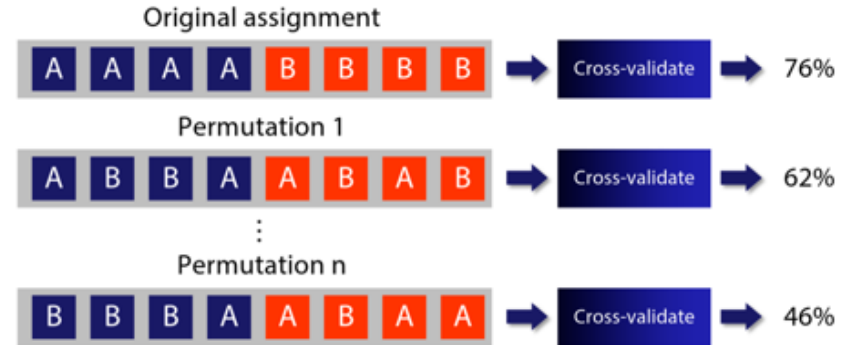
1. Calculate statistic (e.g. accuracy) using normal procedure
2. Permute labels and repeat same procedure
3. Repeat n times or until exhaustion

For exhaustive test: p -value is $\frac{k}{n}$ where k is number of permutations with equal or higher accuracy and n is number of all permutations (includes original result)

For non-exhaustive test (aka Monte-Carlo permutation test): $\frac{k+1}{n+1}$



Typical procedure: Label permutation (i.e. random shuffling)



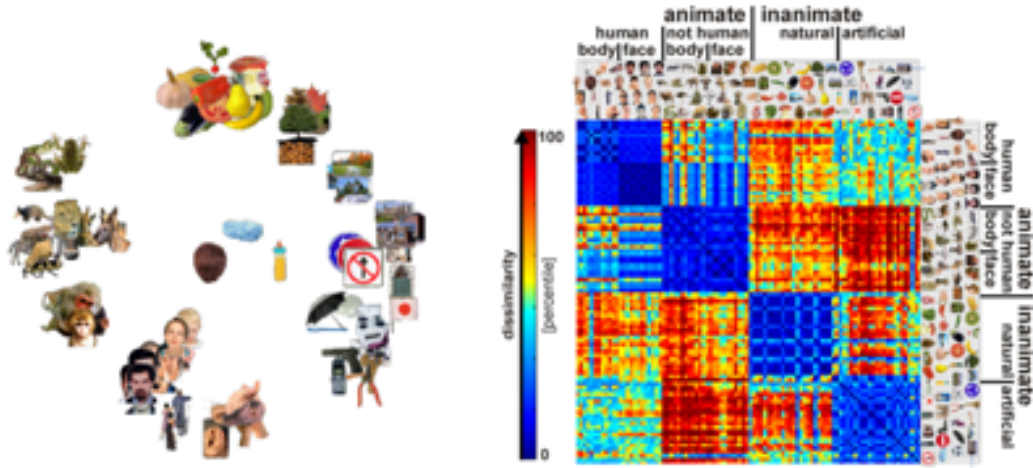
p -value: How often 76 % reached or exceeded by n permutations?

■ ■ Original labels A B Current labels

Step 6: Statistics - confusion matrix

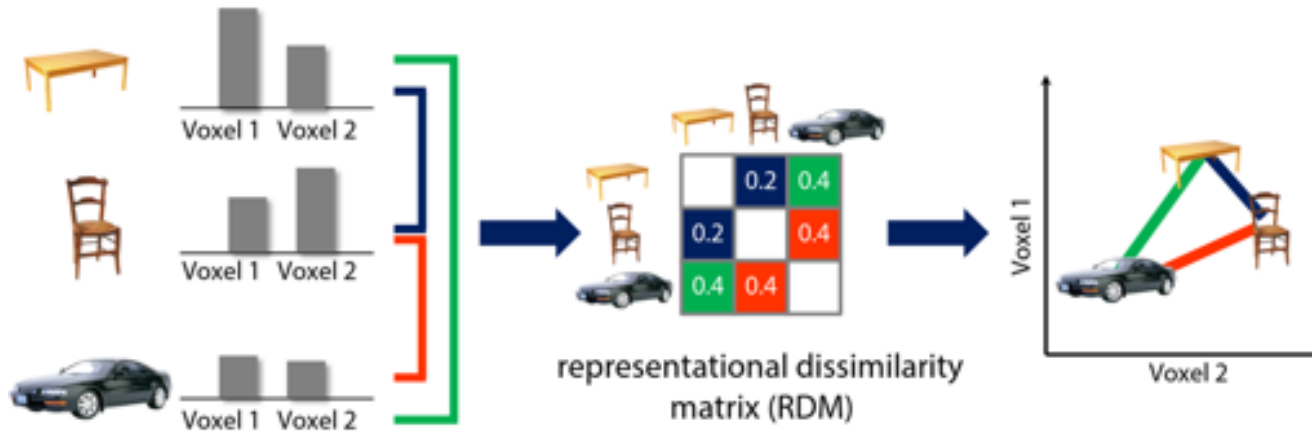
Step 6: Statistics - representational similarity analysis

A multivariate pattern analysis method to investigate the *content* and *format* of representations



Step 6: Statistics - representational similarity analysis

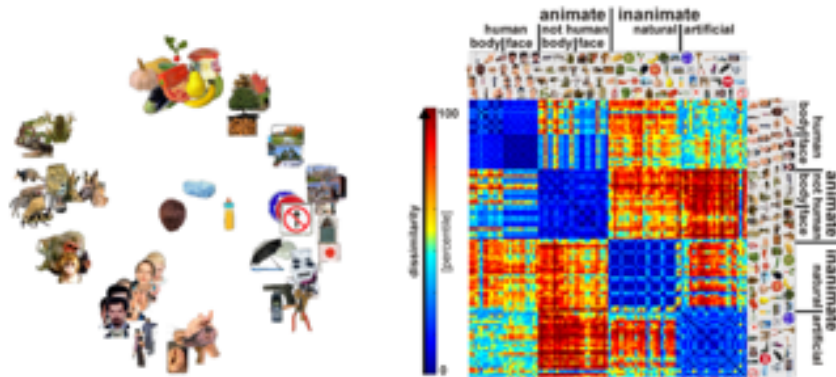
- In RSA, we take our multivariate patterns (e.g. voxels) and calculate pairwise dissimilarities (e.g. Euclidean distance or $1 - \text{Pearson's } r$)



Step 6: Statistics - representational similarity analysis

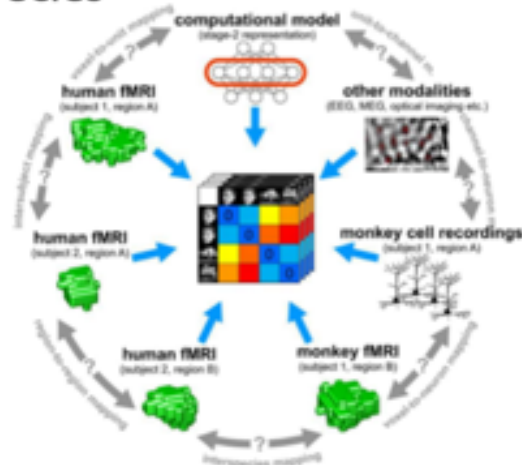
1. Simple exploratory approach to characterize **multidimensional representations**

Example: How does human IT represent object categories?



Step 6: Statistics - representational similarity analysis

2. Representational (dis)similarity matrices can be seen as a **common language to study representations** across methods (MEG, fMRI, cell recording, ...), brain regions, humans and species

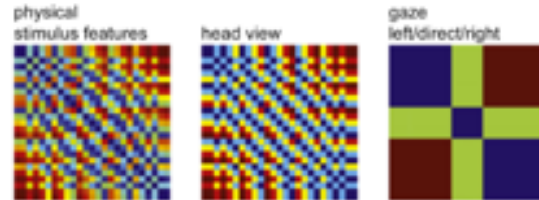
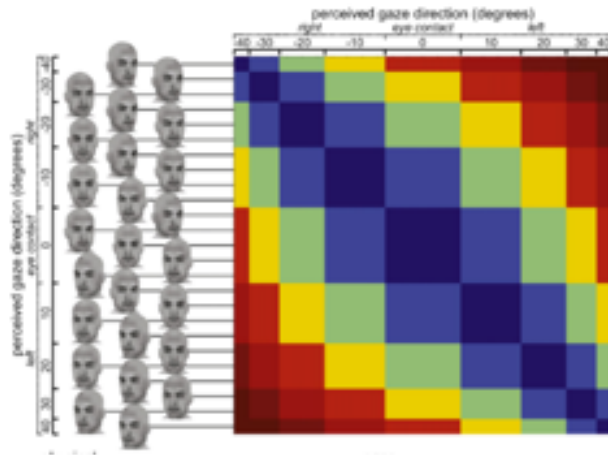


Kriegeskorte et al (2008) – Frontiers; Kriegeskorte & Kievit (2013) – TICS

Step 6: Statistics - representational similarity analysis

3. Representational similarities can be used for **testing models of cognition**

Example: Which facial features does a brain region represent?



More resources

Experimental design, methodological choices etc:

<https://fmrif.nimh.nih.gov/public/other-courses/mvpa>

Conceptual lectures by Rebecca Saxe:

<https://cbmm.mit.edu/fmri-bootcamp>

Tutorials:

<https://brainiak.org/tutorials/>