## Data mining



### What do you do with LOTS of data?

#### General guidelines:

1. Look at your data

perform any/all preprocessing steps use many visualizations, especially of raw, unaveraged data

- scatterplots, heatplots, histograms

look at distributions, identify outliers

test hypotheses with encoding & decoding approaches (confusion matrices!)

look for trends and patterns (unsupervised methods)

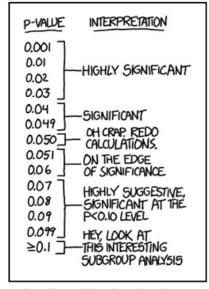
- dimensionality reduction, clustering

## What do you do with LOTS of data?

#### General guidelines:

- 1. Look at your data
- 2. But don't look at all of your data

## Inflation bias, a.k.a. "p-hacking"



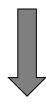


https://imgs.xkcd.com/comics/p\_values.png

## What do you do with LOTS of data?

#### General guidelines:

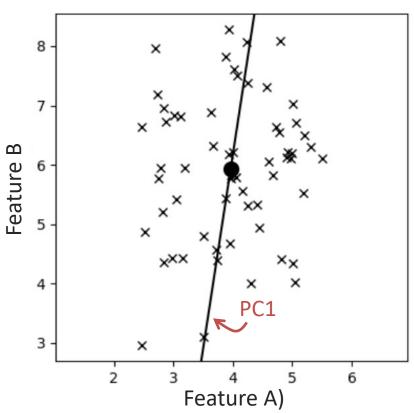
- 1. Look at your data
- 2. But don't look at all of your data



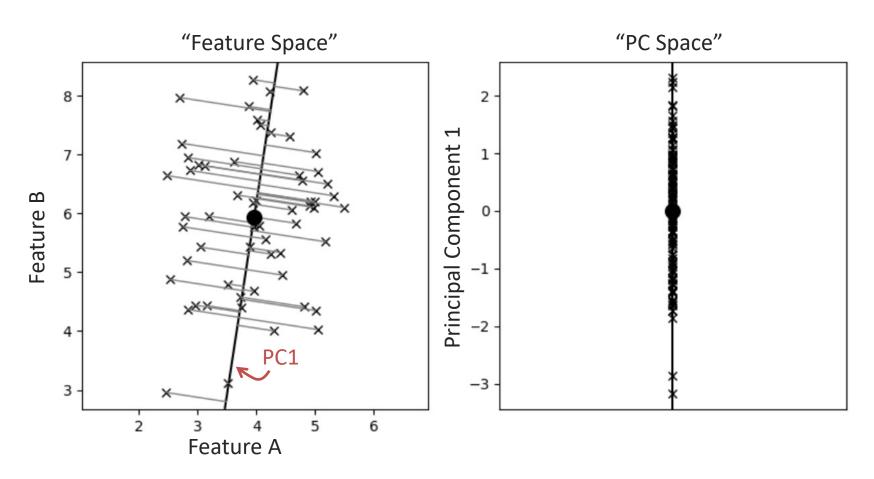
## Best practices to avoid p-hacking

- Create your own replication study by mining half your data and holding half out for test
- 2. Use visualizations to see patterns, do stats only at the end

## Principal component analysis (PCA)



## Principal component analysis (PCA)



#### Let's create some fake data to see how PCA works:

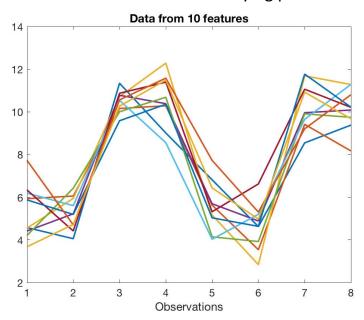
10 features (e.g. neurons) 8 observations (e.g. a measure in 8 different conditions) All features reflect an underlying pattern:

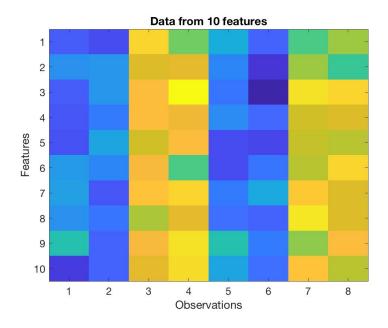
```
% Let's make some features that follow an underlying patterns
% We'll initialize the pattern as responses observed to 8 differnt
% conditions (of in 8 different observations)
pattern1 = [5 5 10 10 5 5 10 10];
var = 1;

% Next we'll make a population of 10 features that follow this pattern with
% some noise
pop1 = [];
% let's make 10 features per subpopulation
for k=1:10
    for j=1:8 %there are 8 observations
        noise = normrnd(0,var); %this will add random noise, drawn from a gaussian centered at 0 with standard deviation = var,
        pop1(k,j) = pattern1(j)+noise;
end
end
```

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#### In matlab:

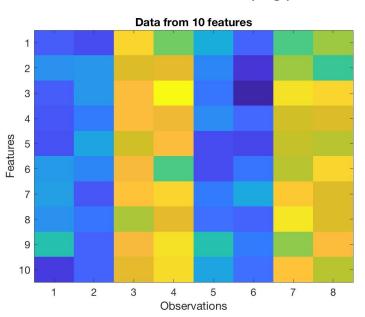
% Now let's use PCA to see how it recovers patterns
[coeff,pcs,~,~,explained]=pca(pop1'); % note that the function that computes PCA has to take input matrices in the correct orientat

#### In R:

pop1 pca <- prcomp(pop1, center = FALSE) # note by default matlab "centers" data R does not # so scale of values differs between matlab and R

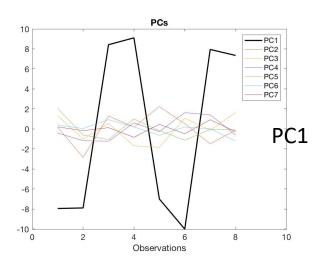
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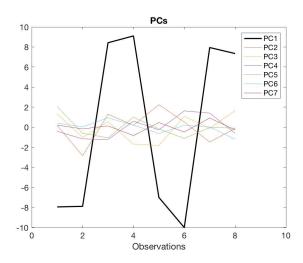
#### PCA can give you:

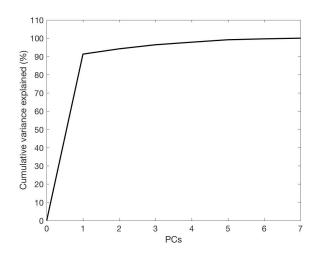
- PCs
- % variance explained by each PC
- feature loadings or "weights" for each PC



#### PCA can give you:

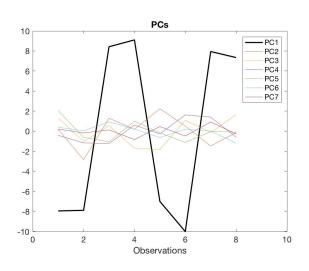
- PCs
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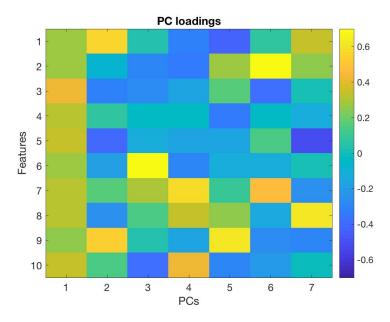




#### PCA can give you:

- PCs
- % variance explained by each PC
- feature loadings for each PC





- PC1 has high weights on each feature
- Other PCs weigh on only 1 or 2 features

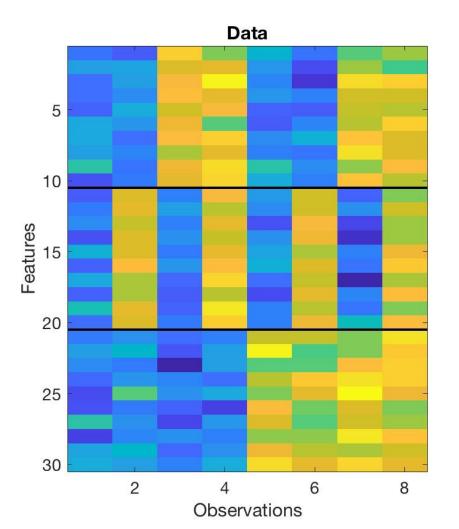
#### Let's add data....

```
30 features (e.g. neurons)
8 observations (e.g. measure in 8 conditions)
3 underlying patterns
```

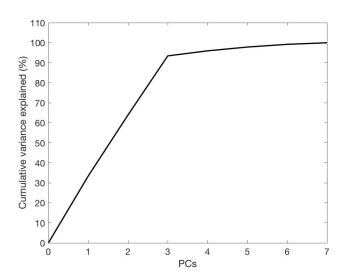
```
% Let's do it again with a more complex data set
% Additional patterns
pattern2 = [5 10 5 10 5 10 5 10];
pattern3 = [5 5 5 5 10 10 10 10];

% and create two more populations that follow different patterns
pop2 = []; pop3 = [];
for k = 1:10
    for j = 1:8
        noise = normrnd(0,var); % noise should be independent for this simulation
        pop2(k,j) = pattern2(j)+noise;
        noise = normrnd(0,var);
        pop3(k,j) = pattern3(j) + noise;
    end
end
pop = [pop1;pop2;pop3]; % Our full feature matrix is all of these subpopulations together
```

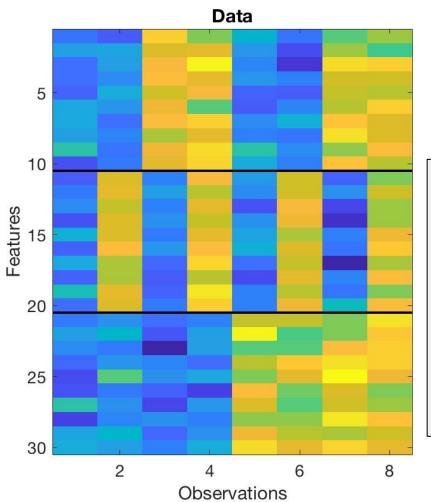
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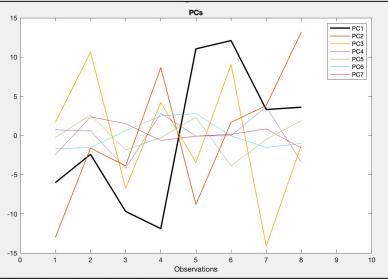
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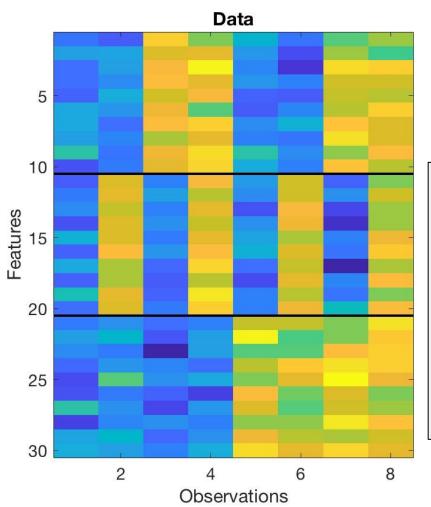
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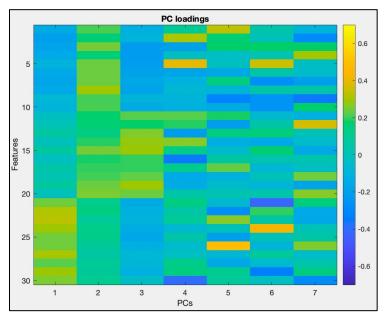


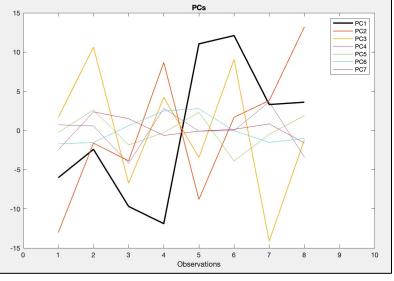
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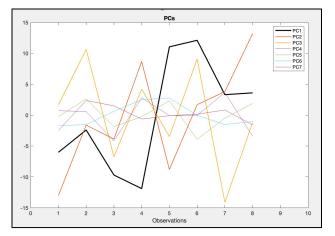


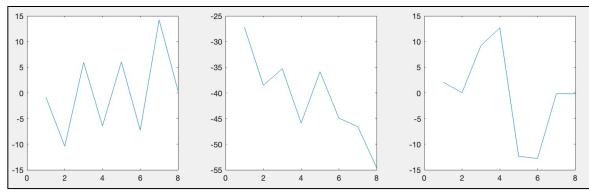


Many algorithms

**PCA** 

Singular value decomposition (SVD – closely related to PCA)
Independent component analysis (ICA) - similar to PCA, but finds
components that have maximal statistical independence





Many algorithms

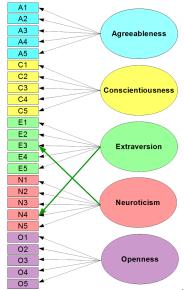
**PCA** 

Singular value decomposition (SVD – closely related to PCA)

Independent component analysis (ICA)

Factor analysis – reduces a large number of variables to a smaller number of "factors" or latent variables by finding maximum

common variance



Many algorithms

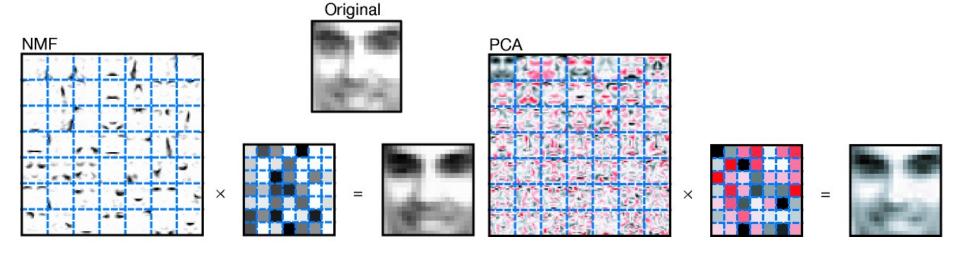
**PCA** 

Singular value decomposition (SVD – closely related to PCA)

Independent component analysis (ICA)

Factor analysis

Non-negative matrix factorization (NMF) – like PCA but constrains weights to be non-negative -> more interpretable in some instances



#### Many algorithms

**PCA** 

Singular value decomposition (SVD – closely related to PCA)

Independent component analysis (ICA)

Factor analysis

Non-negative matrix factorization (NMF)

demixed PCA (dPCA)

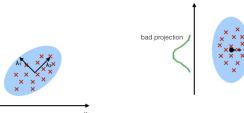
Linear discriminant analysis (LDA)

Bespoke statespace analyses

Methods that aim to find interpretable components

#### PCA:

component axes that maximize the variance



#### LDA:

maximizing the component axes for class-separation

good projection: separates classes we

#### Many algorithms

**PCA** 

Singular value decomposition (SVD – closely related to PCA)

Independent component analysis (ICA)

Factor analysis

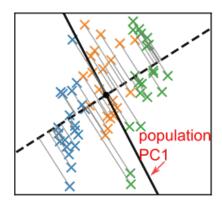
Non-negative matrix factorization (NMF)

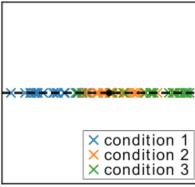
demixed PCA (dPCA)

Linear discriminant analysis (LDA)

Bespoke statespace analyses

Methods that aim to find interpretable components





#### Homework #9

(first part)

#### HW9: Data mining

You have recorded pupil responses in a subject viewing different images. The data are saved in *data.txt*, which includes 500 trials where each trial is 1200ms long

- 1. Plot the mean pupil response over all trials
- 2. Do PCA across trials (Hint: each PC should be1200 elements long, and there should be 500 of them). How much variance does the first PC account for? How many components account for >=90% of the variance?
- 3. Plot the first principal component.
- 4. Run k-means 10 times with k=2. For each run, plot the cluster centers you obtain.